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# Utilization of near-isogenic lines to identify genes underlying iron-efficiency QTL

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**Utilization of near-isogenic lines to identify genes underlying  
iron-efficiency QTL**

by

**Gregory Allan Peiffer**

A dissertation submitted to the graduate faculty

In partial fulfillment of the requirements for the degree of

DOCTOR OF PHILOSOPHY

Major: Genetics

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2011

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## **Dedication**

To my parents, Michael and Sharon and my brother, Andrew for always supporting and encouraging me to make the best out of every situation and always pushing me to achieve more.

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## **Abstract**

Nutrient deficiencies are a significant abiotic stress of soybean. Iron deficiency chlorosis is a major concern in the upper midwestern region of the United States due to the prevalence of calcareous soils. Soybean's susceptibility to iron stress results in yield losses into the hundreds of millions each year. Understanding the molecular differences between resistant and susceptible cultivars will significantly affect future yield and revenue. Through the use of near-isogenic lines (NILs), molecular markers, and gene expression we have identified the donor parent introgressions through both classical SSR mapping and a novel method of SNP clustering which can be preformed using data generated through either chip-based SNP genotyping platforms or identified de novo through re-sequencing techniques. By aligning the newly constructed introgression map with the previously identified Fe efficiency QTL we identified a region on chromosome 3 where the two were positionally coincident. To further narrow this region of interest, the NIL was backcrossed an additional generation to the recurrent parent in order to identify recombinations within the chromosome 3 introgression. These lines were identified as Sub-NILs. Recombinants were identified in regular intervals throughout the introgression and phenotyped. Donor parent alleles identified within a 250 kb region represented the minimum interval differentiating the efficient and inefficient Sub-NILs. A second NIL sharing the same donor parent was screened for introgressions. The only region of the genome the two NILs shared alleles from the donor parent, introgressions, were

localized to the same region on chromosome 3 further adding support to the importance of these alleles. Eighteen genes were annotated within the region and were screened for gene expression differences in soybean roots 24 hours following the removal of iron in the growth medium. Two of the genes were differentially expressed between sufficient and insufficient iron conditions. Interestingly, these genes are homologs of two transcription factors in *Arabidopsis thaliana* known to function in the iron response pathway. Sanger sequencing of these two genes identified a significant mutation that deletes 4 amino acids in the susceptible lines. We hypothesize that this deletion disrupts the FIT / bHLH heterodimer that has been shown to induce known iron acquisition genes.

## Chapter 1. General Introduction

### Rationale

Iron is an integral nutritional requirement for plants and animals alike (Clarkson and Hanson, 1980; World\_Health\_Organization, 1996; Bogden and Klevay, 2000; Welch and Graham, 2004). The negative effects of iron deficiency are seen across kingdoms. Worldwide iron deficiency anemia is a concern with 30% of the human population suffering, however in developing countries such as India greater than 85% of menstruating or pregnant women suffer from severe anemia (Kapur et al., 2002; Theil, 2004). Iron fortification of foods is a viable technique for counteracting anemia in first world countries. However, the cost of these foods is out of reach for people in developing countries (Kapur et al., 2002). Increasing the iron content in locally grown crops would make a far greater impact on the severity of this disorder.

Before significant impacts on human nutrition can be made, the molecular basis of iron uptake and utilization in plants must first be delineated. Iron deficiency chlorosis (IDC) is caused when plants lack sufficient iron for normal growth (Mori, 1999). IDC is manifested phenotypically by interveinal chlorosis, stunted growth and significant end of season yield losses (Abadía et al., 1999). Though IDC should not be thought of in mere economic terms, in 2004 soybean losses were reported up to 25%, equivalent to 120 million dollar loss (Hansen et al., 2004). In Iowa and Minnesota alone, crop losses were greater than 10 million dollars (Hansen et al., 2004). For

these reasons, many breeding populations have been developed to study underlying mechanisms of IDC.

IDC is a complex trait, as the mechanism seems to be population specific. It has been shown to be the result of a single gene with modifiers or a polygenic trait with many genes contributing to phenotype (Cianzio et al., 1980; Cianzio and Fehr, 1982). The single gene hypothesis is supported by a QTL responsible for >70% of the phenotypic variation (Lin et al., 1997). *Lycopersicum esculentum* and *Arabidopsis thaliana* have been used as model systems for studying the reduction strategy of iron uptake. Using these systems, researchers have identified many of the genes responsible for reduction and transport of iron from the rhizosphere (Brown et al., 1971; Grusak and Welch, 1990; Ling et al., 2002; Vert et al., 2002; Connolly et al., 2003; Bauer et al., 2004; Jakoby et al., 2004; Bauer et al., 2007). Furthermore, through recent advances in protein-protein interactions and gene expression studies, some of the co-regulators of the accepted reduction strategy transcription factors have been identified (Yuan et al., 2007).

Near-isogenic lines (NILs) have been created in soybean to study IDC in a common genetic background (Bernard, 1975). It has been suggested that reduction is the rate-limiting step in the iron acquisition process (Grusak and Welch, 1990; Connolly et al., 2003). O'Rourke and colleagues (2007) hypothesized that regulation of the reductase gene is the most likely difference between the NILs.

## Research Objectives

The purposes of the current study were to: (1) map donor parent introgressions in two iron-inefficient NILs and align the introgressions with previous iron-efficiency QTLs; (2) use further backcrossing to narrow the donor parent introgression in the NILs; and (3) identify genes differentiating the response of the NILs to iron stress through coordinate mapping and gene expression analysis using qRT-PCR and RNA-seq.

## Dissertation Organization

This dissertation is organized into five chapters. The first chapter contains the introduction, research objectives and dissertation organization. The second chapter is a literature review presenting current and previous research in iron deficiency chlorosis. Chapters 3 and 4 are each presented as complete manuscripts. Chapter 3 has been published in Plant Physiology, while chapter 4 has been submitted to Plant Physiology. Chapter 5 contains overall conclusions derived from this research in addition to my recommendations for further research.

Chapter 3 entitled, 'An integrative approach to genomic introgression mapping', was published in 2010 volume 154 of Plant Physiology (Severin, Peiffer, et al., 2010).

The donor parent introgressions were identified using polymorphic SNPs from chip based genotyping platforms or next-generation sequencing. Significant clusters of SNPs were used to identify regions of introgressions. Identified introgressions were

confirmed with SSR marker genotyping between the NILs. Collaboration was an invaluable aspect of the manuscript genesis. Gregory A. Peiffer and Andrew J. Severin conceived the concept of SNP clusters differentiating NILs. Andrew J. Severin, Gregory A. Peiffer and Robert M. Stupar conceived the analysis and co-wrote the manuscript. Gregory A. Peiffer generated plant material for next generation sequencing and performed the SSR marker analysis of the NILs and donor parent. David L. Hyten performed the Illumina GoldenGate analysis; Yung-Tsi Bolon contributed material and Illumina GoldenGate data for the protein NILs. Bruna Bucciarelli and Jamie A. O'Rourke generated the Single Feature Polymorphism (SFP) data. Wayne W. Xu performed the SFP data analysis and RNA-seq data analysis using Method-1. Andrew J. Severin performed the RNA-seq data analysis using Method-2. Andrew J. Severin and David Grant created the bootstrapping method. All authors discussed the results to improve the analysis.

Chapter 4 entitled, 'Identification of candidate genes underlying an iron-efficiency QTL in soybean', has been submitted to Plant Physiology and focused on the major NIL donor parent introgression on chromosome 3. Further backcrossing of the NIL with the recurrent parent and coordinate phenotyping narrowed the donor parent introgression. Candidate genes within the narrowed introgression were then analyzed for gene expression changes in iron sufficient and iron deficient conditions. Two transcription factors were differentially expressed at 24 hours in iron stressed roots. Sanger sequencing identified a significant mutation in one of the transcription



factors identified as a homolog of a major regulator gene in the iron response pathway. Gregory A. Peiffer wrote the manuscript, conceived the hypothesis and experimental design through the introgression identification, narrowing and gene expression changes. Keith E. King and Nicholas C. Lauter performed Fe efficiency QTL analysis in the Anoka x A7 population. Andrew J. Severin contributed to the gene expression analysis. Shun Fu Lin provided phenotypic scores of the Anoka x A7. Silvia R. Ciazio generated the population. Randy C. Shoemaker and Silvia R. Ciazio provided guidance and editorial assistance for the manuscript.

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## Chapter 2. Literature Review

Soybean (*Glycine max* (L.) Merr.) is the number one oilseed crop in the world and a major source of protein for animal consumption. Soybean is a multi-purpose commodity used for everything from a simple appetizer to the production of ink and plastics. Production of a large enough quantity of bean to meet the demand requires a healthy crop. In the last decade, United States annual soybean production averaged of 2.98 billion bushels (USDA-NASS). The majority of this crop is produced in the Midwestern United States, which produced a total of 2.85 billion bushels or 86% of the annual yield in 2010 (USDA-NASS). The soils in these states are often calcareous, meaning they have a high calcium carbonate level and a pH around 8 (Inskeep and Bloom, 1987). The high pH and high calcium carbonate levels typically result in a nutritional disorder known as iron deficiency chlorosis (IDC).

### Iron deficiency chlorosis

Iron deficiency chlorosis manifests itself as an interveinal yellowing of the leaves with the persistence of green veins unless the deficiency is severe (Bienfait, 1986). The loss of pigment is due to the plants' inability to manufacture chlorophyll. Iron is an immobile element in the plant, meaning that chlorosis symptoms will only be observed in young leaves and are not consistently expressed throughout the plant. Therefore, chlorosis severity scores are determined from a particular leaf, and are

scored on a continuous five-point visual scale (Cianzio, 1979; Froehlich and Fehr, 1981). A score of one is reserved for a leaf with no yellowing and a score of five is reserved for a completely necrotic and desiccated leaf. A three corresponds with the beginning of interveinal chlorosis and a score of four requires necrotic spots (Cianzio, 1979). The 5-point visual scale has been shown to directly correlate with end of season yield losses. Froelich and Fehr (1981) reported a 20% decrease in yield for each unit increase in the visual score. In a 2003 mail response survey, 99% of the respondents indicated that IDC was a serious issue in their fields and that an estimated 25% of the crop was affected (Hansen et al., 2003). The annual cost to the soybean community due to IDC was calculated at nearly 120 million dollars (Hansen et al., 2004). Today soybean bushels are selling nearly seven dollars higher, increasing the cost to soybean growers to approximately 260 million dollars a year.

Iron deficiency isn't a problem of iron abundance, but rather of plant availability, as iron is the second most abundant metal and the fourth most abundant element in the earth's crust (5.63%) (Taylor, 1964). Iron exists in two biologically relevant states, soluble ferrous iron ( $\text{Fe}^{2+}$ ) and insoluble ferric iron ( $\text{Fe}^{3+}$ ) (Vance, 1994). Soil iron typically exists in the ferric form bound as a part of insoluble oxyhydroxide polymers (Guerinot and Yi, 1994). Free ferrous iron is converted to insoluble ferric complexes very quickly in high pH soils. At a biological pH of 7.0, 90% of the soluble ferrous iron is oxidized within an hour and at a pH of 8.0 the same reaction takes place in less

than 30 seconds (Vance, 1994). The concentration of soluble iron in calcareous soils is approximately  $10^{-10}$  M due to the rapid oxidation and hydrolysis of ferrous iron (Lindsay, 1995). Plants typically require soluble iron at a concentration of  $10^{-4}$  M to  $10^{-9}$  M (Guerinot and Yi, 1994; Lindsay, 1995). Plants have devised multiple mechanisms to overcome the issue of iron availability, without which most could not survive in soils with pH greater than 5.0 (Lindsay, 1995).

Iron forms complexes with a multitude of ligands and can alter its redox potential depending on its conformation (Hell and Stephan, 2003). Its variable redox potential makes iron a prime candidate for electron transport in respiration and photosynthesis. Iron-sulfur clusters are essential for photosynthesis and many other plant pathways (Holm et al., 1996; Staples et al., 1996; Dai et al., 2000). Iron-sulfur clusters are most common in the  $\text{Fe}_4\text{S}_4$  or the  $\text{Fe}_2\text{S}_2$  varieties (Flint and Allen, 1996) although they exist in other ratios (Holm et al., 1996). Iron-sulfur clusters are used to transfer electrons through photosystem I, photosystem II, and the cytochrome b/f complex during the photosynthetic light reactions (Briat et al., 2007). The dark reactions are made possible through the use of iron-sulfur clusters via ferredoxin:thioredoxin reductase, which reduces thioredoxin and in turn activates ribulose 5-phosphate kinase, sedoheptulose 1,7-bisphosphate and fructose 1,6-bisphosphate (Staples et al., 1996; Imsande, 1998; Dai et al., 2000). The above reactions illustrate only a couple of the important roles that iron plays in plant growth

and development. Other roles include nitrate assimilation, sulfur assimilation, and TCA cycle control (Imsande, 1998).

While iron has many benefits for plant growth and development, it requires tight control within the plant. Iron's redox potential, when uncontrolled, has detrimental toxic effects (Hell and Stephan, 2003). Reactive hydroxyl radicals ( $\text{OH}^\bullet$ ) are formed when free iron reacts with superoxide or hydrogen peroxide through a process called the Fenton reaction (Briat, 2002). Hydroxyl radicals cause rapid, nonspecific oxidation of biologically significant compounds that can result in severe damage or death (Graf et al., 1984). Thus, iron within the plant is tightly regulated and shielded from oxygen (Hell and Stephan, 2003). Newly acquired iron is quickly bound by nicotianamine (NA) as the Fe-NA complex makes a poor Fenton reagent (Von Wirén et al., 1999; Hell and Stephan, 2003).

### Strategy I and Strategy II response

Two strategies for iron uptake have been identified in plants (Marschner et al., 1986; Römheld, 1987). The first includes all dicot and non-graminaceous plants such as, *Arabidopsis thaliana*, *Lycopersicum esculentum*, *Pisum sativum* and *Glycine max*. Plants in this group respond to iron stress by acidifying the rhizosphere through the release of hydrogen ions across the root membrane. The lower soil pH increases  $\text{Fe}^{3+}$  solubility in the soil, freeing it from complex soil mineral particles. Soluble  $\text{Fe}^{3+}$  undergoes reduction to  $\text{Fe}^{2+}$  at the plasma membrane via an inducible  $\text{Fe}^{3+}$

reductase (FRO) (Brown, 1978). The reduced Fe is then quickly transported into the root via an iron related transporter (IRT) (Brown, 1978). In addition to the external scavenging methods, strategy I roots induce other physiological changes that include an increase in organic acid concentration in the root to chelate Fe to prevent precipitation or free radical generation, root morphology changes as well as root hair development (Schmidt et al., 2000; Hell and Stephan, 2003). Strategy II plants, including wheat, barley, rice, and maize differ from the strategy I plants in that they secrete ferric chelators, known as phytosiderophores (Hell and Stephan, 2003). Phytosiderophores have been defined as “low molecular weight, virtually Fe(III) specific ligands produced as scavenging agents in order to combat low-iron stress” (Neilands and Leong, 1986). They are created via a one step reaction by nicotianamine synthase that binds three molecules of methionine to nicotianamine (Hell and Stephan, 2003), and are released in a diurnal pattern into the soil (Graham and Stangoulis, 2003). In the soil matrix phytosiderophores scavenge and chelate iron in either the  $\text{Fe}^{2+}$  or the  $\text{Fe}^{3+}$  valencies and then import the entire iron bound complex across the root membrane via a specific transporter identified in maize as the yellow stripe 1 (ys1) gene (Curie et al., 2001; Hell and Stephan, 2003).

### Transcription factor activity

Model organisms for the study of strategy I include *A. thaliana*, pea (*P. sativum*) and tomato (*L. esculentum*), though most of the previous work has been generated from *A. thaliana* and *L. esculentum* studies (Grotz and Guerinot, 2006). There has yet to



be a confirmed gene or mechanism that explains the process of sensing iron stress to initiate transcriptional changes. However, nitric oxide has been implicated in multiple studies as transmitting the iron response signal to initiate transcription of iron acquisition genes (Graziano et al., 2002; Graziano and Lamattina, 2005; Arnaud et al., 2006; Graziano and Lamattina, 2007). The first well-characterized transcription factor in the iron response pathway, FER, was identified in *L. esculentum* through reciprocal grafting experiments (Brown et al., 1971; Brown and Ambler, 1974). Brown et al., (1971) observed that for effective iron uptake a functional FER gene was required in the roots but not in the leaves. More recently, the LeFER transcription factor has been cloned and confirmed to function as part of the iron deficiency transcriptional response (Ling et al., 2002). The *A. thaliana* homolog of the LeFER gene, FER-like Iron Deficiency-Induced Transcription Factor (AtFIT) (Bauer et al., 2007), has been identified as At2G28160 (Bauer et al., 2004; Jakoby et al., 2004; Yuan et al., 2005). Although homologs, the expression patterns of LeFER and AtFIT differ. AtFIT is expressed only under iron stress conditions (Colangelo and Guerinot, 2004; Jakoby et al., 2004). In contrast, LeFER is expressed regardless of iron status in the plant (Ling et al., 2002). LeFER and AtFIT function as transcriptional regulators essential for the induction of IRT1 and FRO2 (Ling et al., 2002; Bereczky et al., 2003; Colangelo and Guerinot, 2004; Jakoby et al., 2004).

AtFIT (AtbHLH029) is a member of the basic/helix-loop-helix (bHLH) transcription factor family. The bHLH family is the largest of the *A. thaliana* transcription factor

families with 174 members (Toledo-Ortiz et al., 2003). Using phylogenetic analysis, the family can be broken into 21 distinct sub-families. The N-terminal region, thought to generate DNA binding specificity, predominantly consists of ~15 basic amino acids (Toledo-Ortiz et al., 2003). The C-terminus of the bHLH protein contains the HLH region thought to function as the dimerization domain (Murre et al., 1989; Ferre-D'Amare et al., 1994). The hydrophobic residues, which make up the two helices, dimerize, allowing the basic N-terminus to bind both halves of the signature E-box recognition sequence (Ma et al., 1994; Shimizu et al., 1997; Toledo-Ortiz et al., 2003). The general consensus sequence for the bHLH recognition domain, the e-box, is 5'-CANNTG-3' with the variable N's providing specificity to this large family of transcription factors (Toledo-Ortiz et al., 2003). For further specification and complexity, the family of bHLH genes has a tendency to function as either homo or heterodimers (Ma et al., 1994; Shimizu et al., 1997; Massari and Murre, 2000; Toledo-Ortiz et al., 2003). Not unlike others in the bHLH family, AtFIT has been thought to interact with another bHLH gene(s) to activate downstream iron regulatory genes (Colangelo and Guerinot, 2004; Grotz and Guerinot, 2006). In the pursuit of these interacting factors, AtbHLH038, AtbHLH039 (Vorwieger et al., 2007; Wang et al., 2007), AtbHLH100, and AtbHLH101 (Wang et al., 2007) were identified as strongly induced under iron stress conditions. These four bHLH genes are members of the sub-family 1b (Wang et al., 2007) or sub-family 2 (Toledo-Ortiz et al., 2003) in the bHLH transcription factor family. This sub-family of genes contains two distinct features that separate them from the rest of the family. The four genes share

a three amino acid insertion within the second helix in the HLH domain, and three of these genes contain a specific cysteine residue in the loop region (Vorwieger et al., 2007).

The central transcription factor AtFIT was shown to bind AtbHLH038 and AtbHLH039 through a yeast two-hybrid analysis (Yuan et al., 2008). This was the first link beyond coordinate gene expression between these three bHLH genes. *A. thaliana* plants over-expressing either AtFIT/AtbHLH038 or AtFIT/AtbHLH039 accumulated more iron in their roots and shoots, and were more tolerant of iron stress conditions than with over-expression of AtFIT, AtbHLH038 or AtbHLH039 alone (Yuan et al., 2008). Yuan et al. (2008) further demonstrated the interaction with dual expression of AtFIT/AtbHLH038 and AtFIT/AtbHLH039 in yeast cells with GUS expression under control of AtFRO2 and AtIRT1 promoters. This indicated that the heterodimers AtFIT/AtbHLH038 or AtFIT/AtbHLH039 were direct regulators of AtFRO2 and AtIRT1.

AtbHLH039 expression has also been shown to be influenced by POPEYE (PYE), a bHLH transcription factor (Long et al., 2010). In *pye-1* mutant plants AtbHLH039 was strongly induced both under iron stress and sufficient conditions, indicating PYE plays a role in regulating AtbHLH039 expression (Long et al., 2010). It is expected that many pathways and regulatory elements are in place to control the expression

of FIT and its interacting factors as FIT plays such a major role in the expression of iron deficiency inducible genes (Colangelo and Guerinot, 2004; Long et al., 2010).

### Ferric reductase oxidase (FRO)

Following the acidification of the rhizosphere, iron is still unavailable for uptake and use in the plant until it is released from organic compounds and reduced to the ferrous state via ferric reductase oxidase (Robinson et al., 1999). Previous research has shown that reduction is the rate-limiting step in the iron acquisition process (Grusak and Welch, 1990; Connolly et al., 2003). Two mutants (*frd1-1* and *frd1-3*) have been identified that lack the ability to reduce ferric iron and therefore cannot efficiently uptake iron (Yi and Guerinot, 1996). It was discovered through cloning and sequencing of the *A. thaliana* ferric chelate reductase gene (*AtFRO*) that it was allelic to both *frd1-1* and *frd1-3* (Robinson et al., 1999). The *frd1-1* mutant contained a stop codon in its first exon and the *frd1-3* mutant harbored a substitution from threonine to methionine in the sixth exon (Robinson et al., 1999). *AtFRO2* is a member of the flavocytochrome family that contains seven other FRO genes that all contain a FAD binding site, a NADPH binding site, four histidine residues, and 8-9 transmembrane helices (Mukherjee et al., 2006). Members of this family have been found in *L. esculentum*, yeast, pea and humans (Dancis et al., 1990; Roman et al., 1993; Chanock et al., 1994; Waters et al., 2002; Li et al., 2004), and are expressed in many different tissues including leaves, shoots, and roots (Robinson et al., 1999; Mukherjee et al., 2006). Expression analysis of the flavocytochrome family in *A.*

*thaliana* has shown that AtFRO2 predominantly functions in the root (Mukherjee et al., 2006). Within 24-hours following iron stress, AtFRO2 transcript levels significantly increase to alleviate the stress (Buckhout et al., 2009).

Over-expression of ferric reductase has shown promising results for increased iron deficiency resistance (Connolly et al., 2003; Vasconcelos et al., 2006; Ishimaru et al., 2007). In 2003, Connolly and colleagues over-expressed AtFRO2 using a constitutive 35S promoter. The transformed plants grew significantly better in iron-stressed conditions compared to the wild-type, which confirmed the claim by Grusak (1990) that reduction was the rate-limiting step in iron acquisition. Since this initial study, the yeast ferric iron reductase gene (refe1/372) was transformed into rice (Ishimaru et al., 2007). The transformed plants performed significantly better under iron stress with a nearly 8 fold yield increase compared to the wild-type plants (Ishimaru et al., 2007). More recently, the *A. thaliana* FRO2 gene was transgenically expressed in soybean grown in hydroponic conditions (Vasconcelos et al., 2006). Transgenic plants showed a significant increase in chlorophyll content in addition to a significant gain in biomass, due to iron deficiency compared to wild-type plants (Vasconcelos et al., 2006). These experiments together indicate that manipulation of the ferric chelate reductase gene can have significant effects on plant growth under iron stress conditions.

### Iron regulated transporter (IRT)

Following reduction by AtFRO2, strategy I plants must transport reduced ferrous iron across the root membrane. The *A. thaliana* iron-regulated transporter 1 gene (AtIRT1) has been implicated as a major component of the iron uptake mechanism (Eide et al., 1996). AtIRT1 has been shown to transport Mn, Zn, Cd and Co in addition to Fe (Korshunova et al., 1999), however, irt1 mutant plants will die prior to seed set unless given exogenous amounts of iron (Varotto et al., 2002; Vert et al., 2002). AtIRT1 is a member of the zinc response transporter (ZRT), IRT-like Protein (ZIP) family that contains 25 members (Guerinot, 2000). The family is further broken down into two sub-families, one consisting mainly of plant derived genes and the other consisting of Animalia genes (Guerinot, 2000). Like AtFRO2, the expression of AtIRT1 mRNA and protein also spike in roots 24 hours following iron stress (Buckhout et al., 2009). Interestingly, AtIRT1 is post-transcriptionally regulated (Connolly et al., 2003) to alleviate excess iron transport (Hell and Stephan, 2003). Post-transcriptional control of AtIRT1 was discovered after two observations: AtIRT1 protein levels were reduced within 12 hours following the re-introduction of iron, and over-expression of AtIRT1 only accumulated AtIRT1 protein in iron-limiting conditions (Connolly et al., 2002). The over-expression of AtFIT/AtbHLH038, AtFIT/AtbHLH039, or the over-expression of AtbHLH039 alone released AtIRT1 from post-transcriptional regulation in both iron sufficient and deficient conditions (Yuan et al., 2007). AtbHLH039 is unable to bind and function as a homodimer, indicating that AtbHLH039 and an unknown factor may regulate the post-

transcriptional control of AtIRT1 (Yuan et al., 2007), further illustrating AtbHLH039's importance in iron homeostasis.

### Identification of QTL associated with iron deficiency chlorosis

Increasing productivity of soybean lines in iron stress conditions can be achieved with the addition of exogenous iron (Abadia et al., 2011), although the most economical and durable method to control IDC is through the identification and utilization of resistant genotypes (Longnecker and Welch, 1990; Mortvedt, 1991; Wiersma, 2005). For decades, soybean breeders have teased out the genomic locations of genes responsible for iron use efficiency in various populations. The earliest study of IDC concluded that iron-efficiency was controlled by a single recessive gene with no maternal effects (Weiss, 1943). Following this initial observation, iron-efficiency was identified as a dominant gene with modifiers (Brown and Caldwell, 1967), a single major gene with co-dominant inheritance (Cianzio et al., 1980), and a quantitative trait controlled by additive gene action (Cianzio and Fehr, 1982). In 1997, Lin et al., mapped quantitative trait loci simultaneously in two populations with two years of leaf chlorophyll concentrations in addition to the standard, visual IDC scores. Their report described two populations, one exhibiting polygenic inheritance and the other a single major gene with modifiers. Polygenic inheritance was observed in a Pride x B216 population with multiple QTL mapping to 6 major linkage groups (MLGs), with no QTL contributing > 30% to the observed phenotype. In the other population studied, Anoka x A7, Lin and colleagues

observed a single major gene with modifiers. In this population, a region on MLG N (chromosome 3) contributed 78.7% to the observed phenotype, further strengthening the argument for two separate mechanisms for IDC control, and that are dependent on the population being studied (Cianzio and Fehr, 1982; Lin et al., 1997).

### Hydroponic growth of soybean lines

Field plots are inherently inefficient for iron deficiency chlorosis QTL mapping and evaluation. Studies are limited to geographic regions harboring calcareous soils, a single growing season during the summer, and the relative severity of chlorosis (Jessen et al., 1986; Lin et al., 2000). To alleviate these limitations, a hydroponic growth system was established by Chaney et al. (1989;1992). Using a hydroponic system, Jessen and colleagues (1988) screened eight genotypes with varying responses to iron deficiency and found a 0.98 correlation between visual scores from calcareous soil and hydroponic environments. To further prove that hydroponic plant growth under iron stress conditions would mimic results obtained on calcareous soils, Lin et al. (2000), expanded their previous iron QTL mapping study to include hydroponic growth and analysis in the same populations previously used (Lin et al., 1997). The polygene hypothesis was validated for the A15 x Pride B216 population with multiple QTL in the same locations as the QTL identified in the field study. In the Anoka x A7 population the QTL on MLG N (chromosome 3) was reconfirmed, explaining 82.9% of the phenotypic variation based on visual scores (Lin et al., 2000).



Since these initial proof-of-concept studies in hydroponic QTL mapping, hydroponics became the standard method used in nutrient studies as it allows for removal of specific nutrients without altering others, allowing for control of other potential variables. Manipulation of specific nutrients is extremely important in mapping specific QTL directly associated with the trait of interest rather than spurious QTL from uncontrolled environmental variables. Hydroponics are the standard for microarray and transcriptome profiling studies for nutrient studies including but not limited to potassium, phosphorus, salt, and iron (Kreps et al., 2002; Hammond et al., 2003; Gierth et al., 2005; O'Rourke et al., 2007; O'Rourke et al., 2007; Buckhout et al., 2009; O'Rourke et al., 2009).

### Near isogenic lines

In molecular breeding it is important to work with a specific population that will most efficiently identify genes associated with the trait of interest. Near isogenic lines (NILs) are immortal lines, in the sense that genetic variation is trapped and preserved through future rounds of self-fertilization to allow for multiple experiments with the same population and quasi-identical genotypes (Keurentjes et al., 2007). Near isogenic lines (in principle) vary only in the alleles responsible for a trait of interest, while identical for all other alleles (Keurentjes et al., 2007), proven to be extremely useful in identification of target genes in many crops (Young et al., 1988; Kim et al., 2008).

Near isogenic lines are obtained through crossing a genotype containing an allele of interest (donor parent) to a genotype containing the desired background alleles (recurrent parent) (Keurentjes et al., 2007). Progeny from the initial cross is then backcrossed to the recurrent parent and the resulting progeny screened for the trait of interest. Only those retaining alleles for the trait of interest are harvested and again backcrossed to the recurrent parent. For each generation of backcrossing to the recurrent parent, the proportion of the donor parent alleles decreases by 50%, meaning the ratio of the recurrent parent alleles to donor parent alleles in the BC1 will be 75:25, BC2: 87.5:12.5, ..., BC6: 99.22:0.78, etc (Fehr, 1987). The alleles introgressed into the recurrent parent can be mapped using molecular markers to identify genomic locations of controlling alleles. After identification of the introgression, greater resolution can be achieved through additional reduction of the donor parent's genetic contribution (Keurentjes et al., 2007) via additional recombination with the recurrent parent (Goodstal et al., 2005). Marker assisted selection is used to identify NILs containing a subset of the whole introgression, called Sub-NILs (Frisch et al., 1999; Goodstal et al., 2005; Kim et al., 2008). Phenotyping of the Sub-NILs can aid in identification of specific genomic regions harboring genes contributing to the trait of interest (Monforte et al., 2001; Brouwer and St. Clair, 2004; Goodstal et al., 2005; Nichols et al., 2006; Kim et al., 2008).

In the soybean community many NILs have been created for public use. The USDA-ARS has specifically created NILs in two separate backgrounds to study iron

deficiency chlorosis (Bernard, 1975; Bernard et al., 1991). Each of these NILs (Clark/PI 547430 and Harosoy/PI 547698) contain introgressed alleles from the iron-inefficient donor parent T203 (Bernard, 1975). The authors were unable to find published literature utilizing the Harosoy / PI 547698 NIL for use in iron studies. The Clark / PI547430 NIL has become the standard NIL for use in studying iron deficiency chlorosis (O'Rourke et al., 2007; O'Rourke et al., 2007; O'Rourke et al., 2009; Rogers et al., 2009).

### Gene expression studies

Genetic research often aims to identify the gene(s) underlying traits of interest. Gene expression analyses using techniques such as microarray analysis or RNA-Seq takes advantage of the transition state between transcription and translation. Following transcription, a gene product exists as messenger RNA (mRNA) that will further be translated into a functional protein. Researchers are able to assess the relative amounts and changes of mRNA levels for a corresponding gene through various iron conditions to better delineate gene(s) associated with stress responses. Gene expression analysis has undergone many different variants since its conception.

The simplest method for tracking gene expression changes for a single gene or a subset of genes is through quantitative reverse transcriptase polymerase chain reaction (qRT-PCR). qRT-PCR is a variation of the PCR, reaction first introduced in

1986 (Mullis et al., 1986) designed to amplify DNA. Through the use of a retroviral enzyme, reverse transcriptase, the reaction produces a complementary DNA copy (cDNA) of the mRNA for further amplification through PCR (Rappolee et al., 1988). This process was first used to quantify cellular mRNA levels in the late 1980's and is still an accepted form of mRNA quantification (Becker-Andre and Hahlbrock, 1989; Wang et al., 1989; Gilliland et al., 1990). qRT-PCR has its limitations, which include high inter- and intra-reaction error, in addition to time and resource inputs required for simultaneous quantification of many genes (Freeman et al., 1999).

With qRT-PCR, research is directed at a list of previously identified candidate genes. This presumption carries with it the very real potential of limiting a researcher's view of broad gene expression changes following a given stress. Microarray experiments have the potential to screen thousands of genes simultaneously, greatly broadening the scope of an experiment (Lockhart and Winzeler, 2000). However, due to the hybridization-based assay it is typically difficult to distinguish between and among members of a gene family. Affymetrix GeneChips® were used in much of the early transcriptome analysis of iron deficiency chlorosis in both soybean and *A. thaliana* (Thimm et al., 2001; O'Rourke et al., 2007; O'Rourke et al., 2007; Buckhout et al., 2009; O'Rourke et al., 2009).

O'Rourke and colleagues identified candidate genes through the use of microarray analysis in soybean grown for ~14 days in iron limiting hydroponic conditions

(O'Rourke et al., 2007; O'Rourke et al., 2007). This initial research was performed using the near isogenic lines Clark and PI547430. PI547430 lacks FRO2 (ferric chelate reductase) activity in roots when to iron-limiting conditions (O'Rourke et al., 2007). Further genomic dissection of the differences between these NILs identified 835 differentially expressed (DE) genes between iron-sufficient and iron-limiting conditions in Clark, and 200 DE genes in PI547430 with only a portion of those genes co-locating to known iron-efficiency QTL (O'Rourke et al., 2009). Considering the observed genomic locations of DE genes, the researchers hypothesized that the genetic difference between these NILs was most likely a mutation within a transcription factor acting transiently to control the expression of the many iron response genes outside known QTL regions (O'Rourke et al., 2009).

In contrast to the early soybean gene expression studies performed at 14 days after iron stress, *A. thaliana* researchers investigated iron deficiency response and gene expression changes at 24 hours post-iron stress (Thimm et al., 2001; Wang et al., 2002; Buckhout et al., 2009). This earlier view into gene expression changes has the potential to identify transcription factor activity responding to the plant's perception of iron stress rather than as a product of that response in the subsequent days. These early *A. thaliana* studies are responsible for the identification of the integral roles of the bHLH subfamily 1b, AtbHLH038, AtbHLH039, etc. (Vorwieger et al., 2007).

Further advancements in sequencing and transcriptional analysis yielded a paradigm shift in the field of transcriptomics (Wang et al., 2009). Next-generation sequencing (NGS) allows for deep sequencing of the transcriptome with higher throughput and improved cost-effectiveness than classical Sanger sequencing (Marguerat and Bähler, 2010). Just as the microarray world was dominated by commercial platforms, the RNA-seq technology is dominated by the Roche 454 system, the AB SOLiD, and the Illumina Genome Analyzer. While they vary as to how sequence reads are processed and produced, they all produce terabytes of sequence information in a few hours of run time. Applications for next-generation sequencing spans from full transcriptome profiling (RNA-seq) to genomic DNA sequencing, both *de novo* and re-sequencing. NGS varies from Sanger sequencing as the new platforms produce short (36 bp – 150 bp) fragments compared to the hundreds of base pairs produced from a single Sanger read. The advantage of the “next-generation” sequence is that tens of millions of reads are produced though a single run, allowing for greater coverage depth (Marguerat and Bähler, 2010). The greater depth of coverage allows researchers to address questions that weren’t possible using older technologies (Ozsolak and Milos, 2011). Microarray studies suffer from a severe disadvantage, the inability to identify gene expression changes for genes expressed in very low or very high levels, RNA-seq can overcome these issues and provide an accurate estimation of expression level (Wang et al., 2009). RNA-seq was recently performed on a soybean line using 13 different tissues ranging from young leaf to nodule (Severin et al., 2010). The data from this experiment allows for

the identification of tissue- and time-specific gene expression and is publically available via various public databases such as Soybase.org.

RNA-seq data can be used for far more than identification of gene expression differences. Post-transcriptional regulation (Marguerat and Bähler, 2010), mapping of intron / exon boundaries (Wang et al., 2009), recognition of alternative splicing patterns and small RNA profiling (Ozsolak and Milos, 2011) are just some of the ways to use the tremendous amount of RNA-seq data generated from a single experiment. The ability to detect single nucleotide polymorphisms is another advantage and possible use of RNA-seq (Cánovas et al., 2010). SNPs have been successfully used to map QTL and also for the identification of NIL introgressions (Hyten et al., 2008; Kaczorowski et al., 2008; Bolon et al., 2010). The SNPs generated from and RNA-seq experiments are by nature located in the coding regions of the gene, and therefore have a greater potential to alter gene function (Cánovas et al., 2010).

### Experimental questions

Fortunately, many researchers have delineated the multiple methods of genetic control and identified the strategy 1 iron response genes and their interactions. IDC gene expression studies have adopted NILs as model populations for such research. Through previous studies, researchers have hypothesized that the different iron stress responses may be due to transcription factors, a mutation in the ferric

reductase gene, or both (O'Rourke et al., 2007; O'Rourke et al., 2009). With the recent release of the Williams 82 soybean sequence and gene annotations (Schmutz et al., 2010) we can now attempt to identify specific genes underlying the iron stress response. To better understand the mechanism behind IDC in soybean we must first identify the mechanism underlying our model system. Identification of the donor parent introgressions in the NILs is paramount to localizing regions of the genome that control iron-efficiency response. Do the donor parent introgressions co-localize with previously identified QTL? Do alleles within the introgressions and QTL vary between the recurrent parent and inefficient NIL? Using modern gene expression analysis, can gene(s) within the region regulated by iron availability be identified? Answers to these questions will significantly impact the soybean research community and provide a better understanding on how to properly use the NILs in further research. The better we are able to utilize the available tools, the quicker we will alleviate yield losses due to this devastating abiotic stress.



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### Chapter 3. An integrative approach to genomic introgression mapping

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## Abstract

Near-Isogenic Lines (NILs) are valuable genetic resources for many crop species, including soybean (*Glycine max*). The development of new molecular platforms promises to accelerate the mapping of genetic introgressions in these materials. Here, we compare some existing and emerging methodologies for genetic introgression mapping: single-feature polymorphism analysis, Illumina Goldengate SNP genotyping, and *de novo* SNP discovery via RNA-Seq analysis of next-generation sequence data. We used these methods to map the introgressed regions in an iron-inefficient soybean NIL and found that the three mapping approaches are complementary when utilized in combination. The comparative RNA-Seq approach offers several additional advantages, including the greatest mapping resolution, marker depth and *de novo* marker utility for downstream fine-mapping analysis. We applied the comparative RNA-Seq method to map genetic introgressions in an additional pair of NILs, exhibiting differential seed protein content. Furthermore, we attempted to optimize the comparative RNA-Seq approach by assessing the impact of sequence depth, SNP identification methodology and post-hoc analyses on SNP discovery rates. We conclude that the comparative RNA-Seq approach can be optimized with sufficient sampling and by utilizing a post-hoc correction accounting for gene density variation that controls for false-discoveries.

## Introduction

Near-Isogenic Lines (NILs) are valuable genetic resources for the identification of genomic regions and alleles responsible for trait variation. This is particularly true within the soybean (*Glycine max*) community, where NILs can be utilized to map the genomic regions responsible for the phenotypic variation of numerous traits, including seed composition, nutrient deficiency tolerance, maturity and several others (Bernard et al., 1991).

Historically, the mapping of NIL introgression sites has relied on a wide range of electrophoresis-based molecular tools, including isozyme, RFLP, AFLP and SSR analyses (Muehlbauer et al., 1989; Muehlbauer et al., 1991; Molnar et al., 2003; Nichols et al., 2006). More recently, automated genotyping technologies have accelerated the efficiency of genetic mapping. Such methods, including Single Feature Polymorphisms (SFP) analysis of microarray data and Single Nucleotide Polymorphism (SNP)-based genotyping methods, have been successfully applied to the mapping of soybean NIL lines and other mapping populations (Hyten et al., 2008; Kaczorowski et al., 2008; Bolon et al., 2010). However, the mapping resolution of all of these platforms is limited by the location and depth of informative markers available for a given species. Additionally, many of the markers will not be polymorphic for the specific set of genotypes utilized in a NIL introgression study.

The recent sequencing of the soybean genome (Schmutz et al., 2010) and recent advances in next-generation sequencing (NGS) technologies have the potential to overcome some of these limitations. Comparative NGS analyses of near-isogenic lines with their respective parental lines offers the possibility of identifying SNP polymorphisms that are unique to each NIL-parent group. Furthermore, comparative NGS analyses offer a potentially greater marker depth than previous mapping methods. Direct RNA sequencing (RNA-Seq) via NGS allows for these goals to be

accomplished at a lower cost, as the sequence coverage per SNP will be enriched within transcribed regions, thereby reducing the total amount of sequence required to confidently identify true polymorphisms.

In this study, we have attempted to map the introgression loci of the soybean NIL IsoClark (PI 547430) relative to its recurrent parent Clark (PI 548533). Previous studies of Clark and IsoClark NIL have characterized the differences between these lines at multiple levels of resolution, including morphological and transcriptional differences (O'Rourke et al., 2007a; O'Rourke et al., 2007b; O'Rourke et al., 2009). Compared to Clark, IsoClark is an iron-inefficient line, putatively caused by the introgression of iron-inefficient genetic material from the donor line T203. Iron deficiency chlorosis (IDC) remains a problem of great economic importance for soybean growers (Hansen et al., 2003). Therefore, the Clark – T203 – IsoClark family represents a soybean NIL family of both scientific and economic importance. Here, we have examined several genotyping technologies to improve the mapping of T203 introgression sites in IsoClark. Furthermore, we have applied our RNA-Seq based methods towards mapping introgression of two additional soybean NILs exhibiting seed composition differences (Nichols et al., 2006). We have compared some of the existing (Affymetrix SFP and Illumina Goldengate) and emerging technologies (Illumina NGS) for soybean introgression mapping, and speculate on what methods and analytical tools will be most useful in the post-genomic era.

## Results

### *Introgression mapping using Affymetrix single feature polymorphisms*

Affymetrix SFP analysis was used to identify putative T203 introgressions in the NIL genotype IsoClark. SFP between Clark and IsoClark were considered indicative of potential T203 introgression sites. We compared 10-day and 14-day root transcripts from Clark and IsoClark, each grown hydroponically in iron-sufficient and iron-limiting

conditions (see Materials and Methods). This analysis identified four obvious SFP clusters in the IsoClark genome, on chromosomes 3, 5, 8 and 16 (Table I, Figure 1). Based on these analyses, it appears that the T203 introgression on chromosome 3 is the largest of the four. Eleven additional SFPs were identified outside of these clusters and were scattered throughout the genome (Figure 1). These SFPs were inferred to be false-positives unless validated by additional genotyping platforms.

#### *Introgression mapping using the Illumina Goldengate platform*

The Illumina Goldengate genotyping platform was used to identify putative T203 introgressions in IsoClark. SNPs between Clark and IsoClark were considered indicative of potential T203 introgression sites. This analysis identified seven loci that were polymorphic between Clark and IsoClark (Table I, Figure 1). Four of these seven loci had been previously identified as likely introgressions based on SFP analysis. One of the remaining loci, on chromosome 13, co-localized with a solo SFP. The two remaining loci, near the top of chromosome 8 and towards the bottom portion of chromosome 4, did not co-localize with any previously identified SFP (Table I, Figure 1).

#### *Introgression mapping using two SNP calling methods on a single-library of Illumina RNA-Seq data*

Illumina RNA-Seq data was used to identify putative T203 introgressions in IsoClark. SNPs identified *de novo* between Clark and IsoClark transcripts were mapped to the soybean reference genome and their genomic positions were considered as potential T203 introgression sites. Altogether, RNA-Seq SNP discovery was performed in four ways: Single-library comparisons using Method 1 (see description below and in “Materials and Methods”), four-library comparisons using Method 1, single-library comparisons using Method 2 (see description below and in “Materials and Methods”), and four-library comparisons using Method 2. This approach

allowed us to compare the sensitivity and accuracy of RNA-Seq SNP discovery across different analytical methods and sequence depths.

For the single-library comparisons, Illumina NGS was performed on the RNA isolated from the 10-day iron-limiting root samples, resulting in 30,897,337 short read sequences. These sequences were then aligned to the *Glycine max* genome (Glyma1.01 genome assembly) to identify SNPs between Clark and IsoClark in protein coding regions. SNPs were considered indicative of T203 genomic introgression sites. Two methods were used to identify SNPs and to gain a measure of confidence in the SNPs determined by each method.

Method 1 used the program SOAP2 (Li et al., 2009b) to align the short read sequences to the soybean genome (Schmutz et al., 2010). SNPs were then identified from the SOAP2 alignment using program SOAPsnp (Li et al., 2009a). Only unique alignments were considered. SNPs were screened for a minimum base-call quality score of ten, average quality score of 20, minimum best hits of four and no ambiguous bases.

Method 2 used the program GSNAP (Wu and Serban, 2010) to align the short read sequences to the soybean genome. GSNAP can handle short read sequences that fall over splice junctions. All mismatches from the best alignment for a read were tallied in a database and a reporting script required the potential SNP to meet the following criteria: a minimum of two unique alignments, average quality score of 20 and a minimum of 80% of the reads uniquely aligned to the position calling the SNP within a sample. This data was further filtered, requiring a minimum coverage of four short read sequences in the NIL pair and requiring a maximum less than 50% in one line calling the SNP within a sample.

There were 172 SNPs identified by Method 1 (Supplemental Table S1) and 255 SNPs identified using Method 2 (Supplemental Table S2) when applied to the 10-day root RNA-Seq single-library comparison. The putative introgression sites previously identified by SFP and Goldengate SNP analyses accounted for 159 of the 172 SNPs identified using Method 1 and 158 of the 255 SNPs identified using Method 2 (Table I). Thus, the larger introgression sites identified using SFP and Goldengate analyses were generally confirmed by the SNPs called by each method, particularly the sites on chromosomes 3, 5 and 16 (Table I). Introgression sites on chromosomes 4 and 13, which were tentatively identified by SFP and/or Goldengate analyses, were strongly confirmed by the RNA-Seq data (Table I). Surprisingly, the two chromosome 8 introgression sites identified by SFP and/or Goldengate analyses were not strongly supported by the RNA-Seq SNP data obtained from this single-library comparison (Table I).

*Introgression mapping using SNP data from multiple Illumina RNA-Seq libraries*

To determine if the quantity of the short read sequence data for identifying introgression sites was limiting sensitivity, we analyzed eight additional Illumina RNA-Seq data sets using Method 1 and Method 2; four from Clark and four from IsoClark plants grown for 19 days after which the plants were exposed to iron sufficient and iron limiting conditions for 24 hours. We refer to this comparison as the ‘four-library comparison.’ The 19 day root and leaf data set contained 91,303,822 short read sequences. Therefore, this experiment included four times the number of experimental conditions and approximately three times the number of short read sequences than was used in the RNA-Seq single-library comparison described in the previous section.

The RNA-Seq four-library comparison of the 19 day samples identified 261 SNPs with Method 1 (Supplemental Table S3) and 469 SNPs with Method 2 (Supplemental Table S4). The Method 1 SNPs appeared primarily in the larger introgression sites,



with only 14 located outside of these regions. The Method 2 SNPs were found outside of the larger introgressions at a substantially higher frequency (Table I); the locations of these SNPs were scattered across the genome. Both Method 1 and Method 2 identified the two introgression sites on chromosome 8 that were essentially missed by the single-library comparison (Table I). However, the Method 2 analysis identified these sites at a much higher frequency.

Without a substantial accumulation of SNPs in one region in the genome or the coincidental overlap of Affymetrix SFPs or Illumina Goldengate polymorphisms with the next generation sequencing SNPs, it may be difficult to distinguish between a site of introgression and an RNA-Seq false positive SNP call. This problem is further confounded by variations in gene density along each chromosome. In order to identify all or nearly all of the prominent T203 introgression sites a statistical method for distinguishing between introgression sites and false positives randomly scattered across the genome was required.

*Accounting for gene density increases the sensitivity of introgression mapping*

The RNA-Seq data identified SNPs based on short read sequences taken from protein coding regions. To account for gene density and to provide a statistical measure of SNP clustering, an algorithm for SNP clustering utilizing a 'bootstrap method' was developed.

The simulated density of SNPs that might be found within a chromosomal interval by random chance was determined by choosing genes at random with replacement. For example, if 204 SNPs were identified on chromosome 3, then the position of 204 genes from chromosome 3 were chosen at random with replacement. The position of the gene was estimated by averaging the start and end coordinates. This process was repeated 1,000 times to obtain an estimate of the mean SNP density and standard deviation for a given interval. Intervals in the genome that contained a

significantly higher density of SNPs than would be expected at random were inferred to be introgressed. An interval was considered to contain a significantly higher density of SNPs if there were three or more SNPs in the interval and the number of SNPs was greater than three standard deviations above the mean SNP density expected by random chance for a given interval. When the bootstrap method was applied to SNPs identified using Method 1 and Method 2 on the RNA-Seq single-library comparison (10 day root), significant intervals were identified on chromosomes 3, 4, 5, 13 and 16 (Figure 2 a,b). SNPs identified using Method 1 and Method 2 from the RNA-Seq four-library comparison (19-day root and leaf) revealed the same introgression sites and additional sites on chromosome 8 (Figure 2 c,d). The introgression sites identified in the bootstrap method are conservative estimates of the full introgression site, but account for 80% of the single-library SNPs, and 93% of the four-library SNPs identified in Table 1.

These data suggest that the quantity and coverage of short read sequences present in the RNA-Seq four-library comparison may alone be sufficient to identify the same introgression sites as was determined from a combination of SFP, Goldengate, and RNA-Seq single-library comparison. More SNPs pass through the filtering criteria with the increased number of reads from the RNA-Seq four-library comparison. Additionally, the sensitivity of the RNA-Seq four-library comparison is aided by the sampling of RNA from different tissue types, assuring that a more comprehensive set of transcripts (and genome space) was surveyed as compared to the single-library comparison.

#### *Application of the advanced NGS introgression mapping on a second NIL pair*

To further validate our method for determined introgression sites, we performed the Method 2 analysis followed by the bootstrapping post-hoc method on an additional set of two NILs, HiPro and LoPro. The two NILs, derived by introgressing *G. soja* into a *G. max* background (see “Materials and Methods”), exhibit differential seed

protein content (Nichols et al., 2006; Bolon et al., 2010). In this case, we were interested in identifying differential introgression patterns between the two lines, therefore the RNA-Seq SNP comparison was performed directly between HiPro and LoPro, rather than the NILs and the *G. max* recurrent parent.

Twenty-eight libraries taken from a variety of tissues and seed developmental stages were included in the RNA-Seq SNP analysis. This data included 97,637,480 short read sequences. Within this seed-protein NIL data, 387 SNPs were identified (Supplemental Table S5). Approximately 40% (153 out of the 387 SNPs) were located within genomic regions determined to be significant based on our bootstrap algorithm. The remaining SNPs in each experiment were randomly scattered across the genome. Our method was able to easily identify the well-known introgressed region on chromosome 20 (Nichols et al., 2006; Bolon et al., 2010). It also identified regions on chromosome 16 and chromosome 18 that were previously unknown (Figure 3). SNP Goldengate analysis on HiPro and LoPro validated all three of these introgressions (data not shown).

#### *Introgression validation*

The IsoClark introgression sites on chromosomes 4, 5, 13, and 16 were confirmed through resequencing by PCR amplification of Clark, IsoClark and T203 DNA (the introgression on chromosome 3 is well-established and did not require further validation). Additionally, candidate introgressions were also validated with SSR markers. SSR markers BARCSOYSSR\_04\_1282, BARCSOYSSR\_04\_1286, BARCSOYSSR\_04\_1297 and BARCSOYSSR\_04\_1299 were polymorphic between Clark and IsoClark on chromosome 4. Similarly, SSR markers Sat\_217 and Sat\_271 were polymorphic on chromosome 5. SSR marker Satt228 was polymorphic on chromosome 8 (nucleotide position 45,272,500). SSR marker Satt490 was polymorphic on chromosome 13. SSR markers BARCSOYSSR\_16\_1047, BARCSOYSSR\_16\_1057, BARCSOYSSR\_16\_1059 and BARCSOYSSR\_16\_1070

were polymorphic on chromosome 16. All markers and positions are available on Soybase (<http://soybase.org>). Only the predicted introgression between 2.0-3.5 Mb on chromosome 8 was not confirmed through resequencing or SSR markers due to problematic primers or lack of SSR markers in that region. This region, however, has additional support from Illumina Goldengate SNP data. A similar introgression validation was performed for the HiPro and LoPro NILs. Resequencing by PCR amplification confirmed the candidate introgression on chromosome 16 but was unable to confirm the introgression on chromosome 18 (the introgression on chromosome 20 is well-established and did not require further validation). However, all three of these introgressions have been validated by Goldengate SNP data (see previous section).

## Discussion

### *Comparison of SFP, SNP Goldengate and NGS RNA-Seq for genetic introgression mapping*

The Affymetrix SFP and Illumina Goldengate SNP methodologies are established as genetic mapping approaches that are far more efficient than electrophoresis-based methods for genome-wide mapping applications (Hyten et al., 2008; Kaczorowski et al., 2008; Bolon et al., 2010). In our introgression mapping for the IsoClark NIL, the SFP and Goldengate platforms primarily identified an overlapping set of putative introgression sites (Figure 1). The Goldengate platform identified seven introgression sites, which were validated in subsequent experiments, indicating that this platform is robust for introgression mapping. The SFP analysis identified five of these seven sites. However, the SFP analysis also identified ten polymorphic markers outside of these larger introgressions, some of which are believed to be false-positives.

The SFP and Goldengate SNP mapping approaches are relatively efficient and inexpensive. However, in our analyses, the Affymetrix SFP and Goldengate platforms identified a relatively small number of polymorphic markers (Table I). The low number of markers limits our ability to resolve the introgression boundaries and leaves open the possibility of missing smaller introgressions altogether.

The RNA-Seq methodologies clearly identified a much greater number of polymorphic loci within the known introgression sites (Table I). The increased marker coverage allowed us to identify the introgression boundaries at a higher resolution. The two introgressions on chromosome 8, however, were exceptional in this regard. The introgression at positions 2.0-3.5 Mb was easily identified by the Goldengate approach and the introgression at positions 43.9-47.0 Mb was easily identified by the SFP approach. It is unclear what properties of the chromosome 8 introgressions cause this phenomenon; the gene content and transcription levels are both relatively high in these regions (Libault et al., 2010; Schmutz et al., 2010). The other five larger introgressions were most clearly identified by the RNA-Seq approach, regardless of which of the four RNA-Seq analyses was considered.

Importantly, the RNA-Seq approach offers two important benefits that standardized mapping platforms do not. First, the SNP markers identified via RNA-Seq are specific to the genetic materials of interest. By contrast, the soybean Goldengate SNP panel is derived from different genetic materials than was used in our study; many of the 1536 SNPs would be non-polymorphic between our original parents, Clark and T203, and would therefore be uninformative for this study. The RNA-Seq data, however, identifies SNPs that are necessarily polymorphic between our genetic materials of interest. The SNPs identified *de novo* by RNA-Seq can be directly used for fine-mapping on subsequent generations of this material using a custom SNP genotyping platform, like the MassARRAY (Sequenom, San Diego, CA) or SNPlex (Applied Biosystems, Foster City, CA) platforms (Ding and Jin, 2009). Second, the

RNA-Seq data may be mined for transcriptional differences or genetic alterations between Clark and IsoClark that may identify candidate genes that drive the differential iron susceptibility observed between the lines. The Affymetrix data will also allow for the analysis of transcript differences, however the RNA-Seq data provides a larger sampling of transcripts and also permits the possible identification of frame-shift or nonsense mutations within introgressed loci.

We noted two primary drawbacks to the RNA-Seq approach. First, this technology is currently more expensive than using standardized platforms. This problem should be mitigated in the near future, as NGS is expected to become more affordable and accessible. Second, this approach targets mRNA transcripts, therefore our marker depth is necessarily biased for gene rich regions. Although we have applied a bootstrapping method to correct for gene density biases, severely gene-poor regions may not be represented in our analyses. Additionally, exonic regions tend to have more highly conserved sequences than non-coding regions. Introgression mapping could be improved if the next-generation sequencing technology were performed directly on DNA rather than RNA. With current technology, this would provide better genomic coverage but may not provide the sequence depth required for confident SNP identification at a reasonable cost. A more cost-effective strategy would be to perform comparative NGS on reduced representation genomic DNA libraries (Van Tassell et al., 2008; Fu et al., 2010). As sequencing technologies improve and the cost per library decrease, the limitations of sequencing depth and read length will no longer be an issue.

Altogether, our data indicate that the RNA-Seq approach offers the greatest depth and resolution for mapping most genomic introgressions, however the SFP and Goldengate approaches were more efficient for mapping certain introgressions. The combination of SFP, Goldengate and RNA-Seq data does not necessarily assure that we have identified all the introgressed loci in these NILs. For example, when we

combined the unique SNPs identified using Method 1 and Method 2, we noted that a cluster of four SNPs was identified within a ~480 kb interval on IsoClark chromosome 2 (positions 42.35-42.83 Mb). Intuitively, it would appear that these SNPs may define a genetic introgression, however this region was not identified as significant by our bootstrap analyses using each Method (1 and 2) individually. Using the methods described here, introgressions greater than 0.5-1.0 Mb can be efficiently mapped with relatively high resolution, assuming there is an adequate level of sequence polymorphism between the parental lines. However, it may be difficult to identify introgressions that are small, located within gene-poor regions, or located within regions of low diversity between parental lines. Identification of such introgressions, such as the putative introgression on IsoClark chromosome 2, may require “manual” rather than automated analytical approaches, along with sufficient validation.

#### *Optimizing NGS RNA-Seq for genetic introgression mapping*

We tested the impact of three different factors on RNA-Seq introgression mapping: 1) Sequence depth; 2) SNP identification methodology; 3) Post-hoc analysis accounting for gene density.

Clearly, the RNA-Seq method is more effective for introgression mapping when the sequence depth and tissue sampling range is expanded. Our data indicates that our four-library comparison with different tissue types and treatments identified greater than 1.5-times more SNPs than a single-library comparison (Table I). Consequently, introgression sites that were either poorly identified or not identified in the single-library analysis (namely, the two introgressions on chromosome 8) were more confidently identified in the four-library comparison.

We applied two different SNP-identification methodologies to the RNA-Seq data, generically called Method 1 and Method 2 (see Materials and Methods). The two

methods were each applied to the RNA-Seq single-library and four-library comparisons of Clark and IsoClark. These two identification methods appeared to offer an interesting trade-off in benefits. Method 1 appeared to be the more conservative approach, as it identified fewer SNPs. However, only 5-7% of the SNPs were located outside of the putative larger introgression regions identified by SFP and Goldengate genotyping; it is unclear what proportion of these SNPs represent false-positive calls. Method 2 appeared to be the more liberal method, identifying far more SNPs than Method 1 (Table I). A high percentage of SNPs fell outside of the putative larger introgressions (~22-38%), indicating that this method may foster a higher rate of false-discoveries. However, Method 2 was more effective at identifying recalcitrant introgressions, primarily the two chromosome 8 introgressions. It is worth noting that the differential SNP discovery rates of Method 1 and Method 2 are not necessarily a function of the algorithms used (Li et al., 2009a, 2009b; Wu and Serban, 2010), but are also influenced by the stringency of the identification parameters. Thus, either method could be performed with greater or reduced stringency, as needed by the user.

The post-hoc bootstrap method was used to distinguish true introgressions from false-discoveries by accounting for regional SNP clustering rates and gene density differences across the genome. This method proved most valuable when applied to the Method 2 SNP calls, as this was the more permissive identification method and presumably identified a higher relative rate of false-positives. The bootstrap method, when applied to the Method 2 four-library comparison SNPs, identified all of the seven larger introgressions, including the recalcitrant introgressions on chromosome 8.

The data analyses presented here covered a range of tissues and conditions and were performed on a well-studied organism with a set of high-quality predicted gene models. Our analyses indicate several regions of introgression that have been



confirmed for two different NILs. However, had the number of expressed genes been significantly lower than what is found in our data sets, it may be prudent when accounting for gene density with our bootstrap method to use as input to our scripts only expressed genes instead of every gene predicted in the genome.

## Conclusion

In this report, we show that SFP, Illumina Goldengate and RNA-Seq are complementary methods for identifying genetic introgressions in near-isogenic lines. We show that the depth of coverage of SNPs identified from next generation sequencing RNA-Seq technology in combination with a bootstrapping method is an effective tool for identifying introgression sites. As new NGS technologies arise (Eid et al., 2009; Rusk, 2009) and become more affordable, NGS of genomic DNA at greater depth will become feasible for mapping purposes.

## Materials and Methods

### *Plant materials*

Two pair of soybean (*Glycine max*) NILs were used in this study: 1) A NIL line selected for differential iron deficiency chlorosis susceptibility and; 2) A NIL pair selected for differential seed protein. The iron-efficient parent line Clark (PI 548533) and the iron-inefficient NIL IsoClark (PI 547430) have been extensively described in previous studies (O'Rourke et al., 2007a; O'Rourke et al., 2007b; O'Rourke et al., 2009). The IsoClark NIL was derived from crossing Clark with iron-inefficient T203 (PI 54619), followed by five subsequent backcrosses to Clark. Subsequent self-mating yielded the iron-inefficient NIL IsoClark.

The seed protein NIL pair were derived from introgressing *G. soja* (PI468916) into *G. max* (A81-356022) and have been previously described (Nichols et al., 2006; Bolon et al., 2010). The BC<sub>5</sub>F<sub>5</sub> plant P-C609-45-2-2 was heterozygous for the LG I protein QTL introgression from *G. soja*. The derived BC<sub>5</sub>F<sub>6</sub> NILs segregated for the LG I protein QTL introgression. The BC<sub>5</sub>F<sub>6</sub> line LD04-15154 (HiPro) maintained the introgression and the corresponding high seed protein phenotype. The BC<sub>5</sub>F<sub>6</sub> line LD04-15146 (LoPro) segregated out the introgression QTL and exhibited the low seed protein phenotype.

*RNA sampling of Clark and IsoClark root tissues from iron sufficient and iron limiting conditions (10-day and 14-day)*

Clark and IsoClark were grown in hydroponic conditions as described in (O'Rourke et al., 2009). Both genotypes were exposed to two different hydroponic treatments, iron sufficient (100  $\mu$ M Fe(NO<sub>3</sub>)<sub>3</sub>) and iron limiting (50  $\mu$ M Fe(NO<sub>3</sub>)<sub>3</sub>). Roots were collected and flash-frozen in liquid nitrogen following ten and 14 days of growth. RNA samples were purified from both Clark and IsoClark root tissues using the TRIzol method (Invitrogen, Carlsbad, CA) and DNAase treated with the Ambion DNA-free kit according to the manufacturer instructions (Applied Biosystems/Ambion, Austin, TX). The samples were then further purified using the RNeasy mini kit (QIAGEN Inc., Valencia, CA). These RNA samples are referred to as the '10-day root' and '14-day root' samples, respectively.

*RNA sampling of Clark and IsoClark tissues following iron-shock (19-day root and leaf)*

Clark, IsoClark and T203 seeds were germinated using germination paper soaked in water for 6 days in an growth chamber set at 27 °C. Plants were grown in hydroponic conditions as described in O'Rourke et al. (2009) in the greenhouse for 13 days which coincided with the fully-open first trifoliate. At this time the plants were placed in either iron sufficient or deficient conditions. Briefly, the plant roots were rinsed in

six buckets of water for 15 seconds minimum in each bucket then returned to a fresh hydroponic bucket either sufficient in iron ( $100 \mu\text{M Fe}(\text{NO}_3)_3 \cdot 9\text{H}_2\text{O}$ ) or deficient in iron ( $50 \mu\text{M Fe}(\text{NO}_3)_3 \cdot 9\text{H}_2\text{O}$ ).

Plants were grown for 24 hours in their new iron environment when the trifoliate, trifoliolate, and roots were harvested and placed in individual tubes and flash frozen in liquid nitrogen until stored at  $-80^\circ\text{C}$ . Total RNA was isolated using RNeasy mini kit (QIAGEN Inc., Valencia, CA) following the Qiagen protocol for everything except the final elution step that was extended by five minutes to optimize RNA concentration. Quality was checked using a NanoDrop Spectrophotometer (Thermo Scientific, Wilmington, DE). These RNA samples from root and leaf in iron sufficient and deficient conditions are referred to as the '19-day' samples.

#### *RNA sampling of the seed-protein NIL*

Seeds from NILs generated from *G. max* (A81-356022) and *G. soja* (PI468916) specific for the LG I seed-protein QTL were grown in growth chambers to mimic Illinois field growing conditions, as described by Bolon et al. (Bolon et al., 2010). Briefly, 14 tissues that included seven stages in seed development were harvested from the two near-isogenic lines: HiPro (LD0-15154) and LoPro (LD0-15146), with a high and low seed-protein phenotype. RNA was extracted as described in Bolon et al. (2010). These RNA samples are referred to as the 'HiPro' and 'LoPro' NIL samples.

#### *Single feature polymorphism analysis*

The 10-day and 14-day root RNA samples were labeled and hybridized to the Affymetrix® GeneChip® Soybean Genome Array according to the manufacturer's instructions (Affymetrix, Santa Clara, CA). Three biological replicates for each genotype by treatment were collected and hybridized.

Single feature polymorphisms (SFPs) between Clark and IsoClark were identified based on the Affymetrix data as previously described (Xu et al., 2009). SFPs between Clark and IsoClark identified at any of the four levels of comparison (10-day iron sufficient, 10-day iron limiting, 14-day iron sufficient or 14-day iron limiting) were included in the downstream SFP analyses. The Affymetrix SFP probes-sets were mapped back to the Williams 82 soybean genome reference sequence (Schmutz et al., 2010). T203 genomic introgressions into IsoClark were inferred based on SFP co-localization clusters.

#### *Illumina Goldengate mapping*

Clark, IsoClark and T203 DNA samples were purified using the Qiagen DNeasy method according to the manufacturer (QIAGEN Inc., Valencia, CA). These DNA samples were genotyped using the Illumina Goldengate 1536-SNP platform for soybean, as has been previously described (Hyten et al., 2010).

#### *Illumina Next-generation sequencing of RNA*

Illumina next generation sequencing platform was used to identify SNPs between Clark and IsoClark. The three Clark and IsoClark RNA biological replicates from the 10-day root tissues in the iron limiting condition were pooled within each genotype and submitted for next generation sequencing analysis. Similarly, the 19-day root and leaf RNA samples grown in iron sufficient and deficient conditions were each pooled among three biological replicates within each genotype and submitted for next generation sequencing analysis. Therefore, eight different pooled samples from the 19-day study were sequenced, consisting of Clark root and leaf in stressed and unstressed conditions and IsoClark root and leaf in stressed and unstressed conditions.

RNA-Seq data acquisition from Illumina sequencing methods were carried out by the National Center for Genome Resources (NCGR). These techniques along with

RNA-Seq data analysis methods for the seed-protein NILs have been previously described in “RNA-Seq Atlas of *Glycine Max*” by Severin et al. (in press). Briefly, poly-A containing RNA isolated from total RNA was converted to cDNA. Illumina adapters were added by ligation and size selected by electrophoresis for ~500 bp fragments. The purified DNA libraries were PCR amplified for 15 cycles and assessed by Nanodrop ND-1000 for quality and quantity before loading onto an Illumina flow cell. Short reads of size 36 bp were obtained and processed through image analysis, base-calling quality filtering, and per base confidence scores. Sequence reads were then aligned to the 8X soybean genome sequence assembly (Glyma1.01 genome assembly). The unique reads obtained from the alignment were normalized using RPKM (Reads/Kb/Million) and used to approximate digital gene expression values for the predicted gene models.

#### *Next-generation sequencing SNP discovery using Method 1*

Software SOAP2 (Li et al., 2009b) and SOAPsnp (Li et al., 2009a) were used for SNP discovery between Clark and IsoClark genotypes using an RNA-Seq single-library comparison (the 10-day iron limiting root samples). A customized pipeline was developed for this analysis. Briefly, 15,260,698 36-base read sequences of Clark and 15,636,639 reads of IsoClark from Illumina sequencing were aligned to the soybean genome sequence (Schmutz et al., 2010) using SOAP2, respectively. Only the unique alignment hits were selected by setting the program parameter  $r=0$ . All position loci of the alignment files were screened by SOAPsnp for SNPs, and pair compared between Clark and IsoClark. The potential SNPs were selected using criteria of minimum base-call quality of ten, average quality of 20, minimum best hits of four, and not an ambiguous base (SNP  $\neq$  “N”).

In order to plot the SNP alignment, all short read sequences that encompassed the SNP positions were extracted from the original Illumina read files. For each SNP, the short reads of Clark and IsoClark, and the 68-base genomic sequences that

encompass the SNPs in the middle were aligned by using emma of EMBOSS suite (Rice et al., 2000). The aligned sequences were plotted using EMBOSS prettyplot program.

The locations of the SNPs discovered were extracted and an R script was created for mapping these SNPs onto soybean chromosomes using a 1000 base window size. The protocol described above is referred to as the RNA-Seq 'Method 1'.

To determine the difference in sensitivity between an RNA-Seq single-library comparison and a four-library comparison, Method 1 was also applied to the four 19 day samples of Clark and IsoClark: root under sufficient iron conditions, root under iron limiting conditions, leaf under iron sufficient conditions and leaf under iron limiting conditions. The pooled 19 day samples contained 32,030,175 36-base read sequences in Clark and 59,273,647 read sequences in IsoClark.

#### *Next-generation sequencing SNP discovery using Method 2*

For comparison, the software GSNAP (Wu and Serban, 2010) was also used for SNP discovery between Clark and IsoClark genotypes on the RNA-Seq single-library comparison (the same 10-day iron limiting root samples as was used for Method 1) and the 19 day four-library comparison. Briefly, Clark and IsoClark reads from Illumina sequencing were each aligned to the soybean genome sequence using GSNAP. The alignment program was set to allow for alignment over a splice junction. Alignments of short read sequences without at least 34 matches were not considered. The following requirements were also needed for a SNP to be called: a minimum of two unique alignments calling the SNP, average base-call quality of 20 and minimum of 80% of the reads uniquely aligned to the position calling the SNP. SNPs were further screened for a minimum of short read coverage of four and a difference in allelic frequency between the NILs of 50%. The protocol described above is referred to as the RNA-Seq 'Method 2'.

### *Statistical significance and visualization of SNP clusters*

A specific number of SNPs were found on each chromosome using Method 1 and Method 2. To determine which regions on the chromosome had a significantly higher density of SNPs than might be found by random chance, the same number of SNPs found on each chromosome was simulated using a bootstrapping protocol (Supplemental File S1). Since the sequence used to identify SNPs was taken from protein coding regions, the locations of the simulated SNPs were generated from the average position of each gene on the chromosome chosen at random. The chromosome was divided into 500,000-nucleotide tandem intervals, resulting in a total of 1908 intervals analyzed across the 20 chromosomes. The average number of simulated SNPs and standard deviation within each 500,000-nucleotide interval were determined from 1000 simulations. SNPs in an interval were considered significant if the number of SNPs was greater than three standard deviations above the simulated SNPs in the interval and the total SNP count in the interval was three or more. Once the intervals with significant SNP clustering were determined, these regions were plotted onto a scaled version of each chromosome using the rectangle drawing function in R. The protocol described above is referred to as the 'bootstrap method'.

### *Laboratory confirmation of genomic introgressions identified in silico*

Genomic regions identified as candidate introgressions were identified and a small portion of the sequence in the region was extracted from Soybase (<http://soybase.org>). Primers were used to PCR amplify Clark, Isoclark and T203 DNA. PCR reactions were conducted using a touchdown method starting with a 60 °C annealing temperature and decreasing by half a degree each cycle for 29 cycles. Choice™ Taq (Denville Scientific, Metuchen, NJ) was used and PCR reactions were at concentrations according to the manufactures protocol.

PCR products were cleaned using an exonuclease 1 and shrimp alkaline phosphatase (EXO/SAP) method. Cleaned PCR products were used in a cycle sequencing reaction. The sequencing protocol was adapted from the Applied Biosystems BigDye® Terminator v3.1 Cycle Sequencing kit (Applied Biosystems, Foster City, CA). Sequencing was done on an Applied Biosystems 3730xl 96-capillary 50 cm array DNA analyzer. Sequence was end trimmed using Applied Biosystems Sequence Analysis v5.2 (Applied Biosystems, Foster City, CA). Sequence ends were trimmed until fewer than four of 20 bases had quality scores less than 20. The sequences generated from each primer pair were aligned using Sequencher v4.9 (Gene Codes Corporation, Ann Arbor, MI).

Additionally, single sequence repeat (SSR) markers were chosen from Soybase (Soybase.org) in candidate regions of introgression in the IsoClark line. PCR reactions were conducted using a touchdown method starting with a 60 °C annealing temperature and decreasing by half a degree each cycle for 29 cycles. Choice™ Taq (Denville Scientific) was used and PCR reaction was at concentrations according to the manufactures protocol. Bromophenol blue loading dye was added to the PCR reaction and loaded onto a 6% polyacrylamide gel ran 250 volts for 2.5 hrs. Bands were visualized at 312 λ using a grayscale digital camera (Scion Corporation). The lowest band was scored and compared to 10 and 100 bp ladder.

### Acknowledgments

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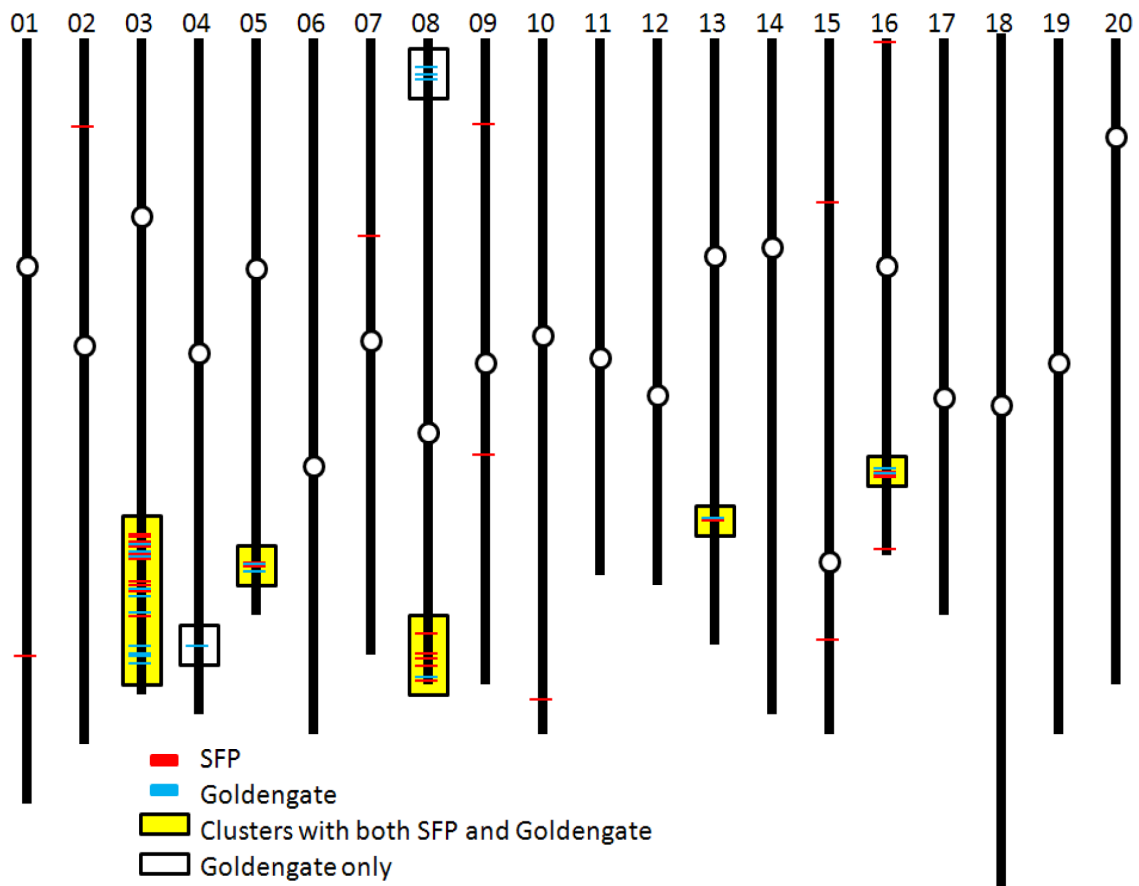
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Table 1. *Introgression mapping comparison*

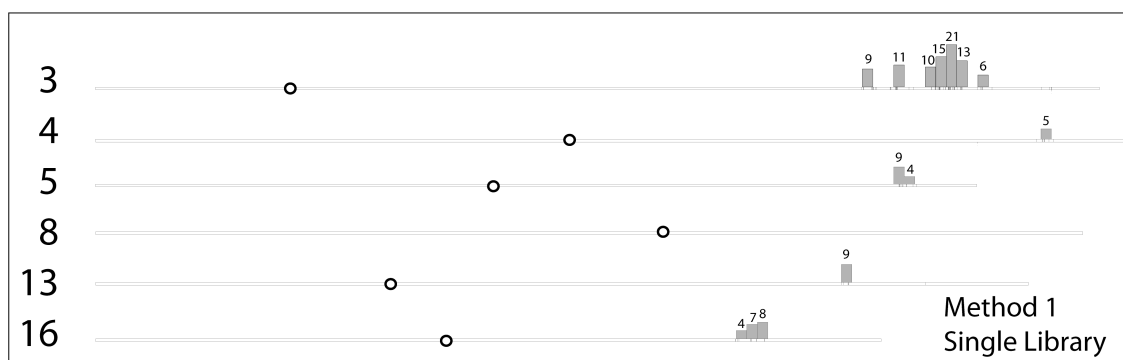
Chromosome	Position	Approximate Size	SFP	Golden Gate	RNA-Seq Single Library (Method 1)	RNA-Seq Four Library (Method 1)	RNA-Seq Single Library (Method 2)	RNA-Seq Four Library (Method 2)
	<i>Mb</i>	<i>Mb</i>						
Gm03	36.3–45.8	9.2	13	15	105	120	102	204
Gm04	44.7–45.6	1	0	1	7	21	6	23
Gm05	38.2–39.5	1	4	2	15	24	17	34
Gm08	2.0–3.5	1.5	0	4	1	2	1	14
Gm08	43.8–47.0	3.2	7	1	0	8	2	17
Gm13	35.5–35.9	0.5	1	1	10	24	7	23
Gm16	30.4–31.9	1.5	6	3	21	48	23	50
Other	N/A	N/A	10	0	13	14	97	104
Total	N/A	N/A	41	27	172	261	255	469

Comparison of Clark-IsoClark polymorphism rates at larger introgression sites using three different genotyping platforms. The number of SNPs identified from the RNA-seq data depends on the SNP-calling algorithm and the number of libraries compared (Single-library comparisons versus four-library comparisons). The RNA-seq method 1 and method 2 analyses protocols are described in the “Materials and Methods.” N/A, Not Applicable.

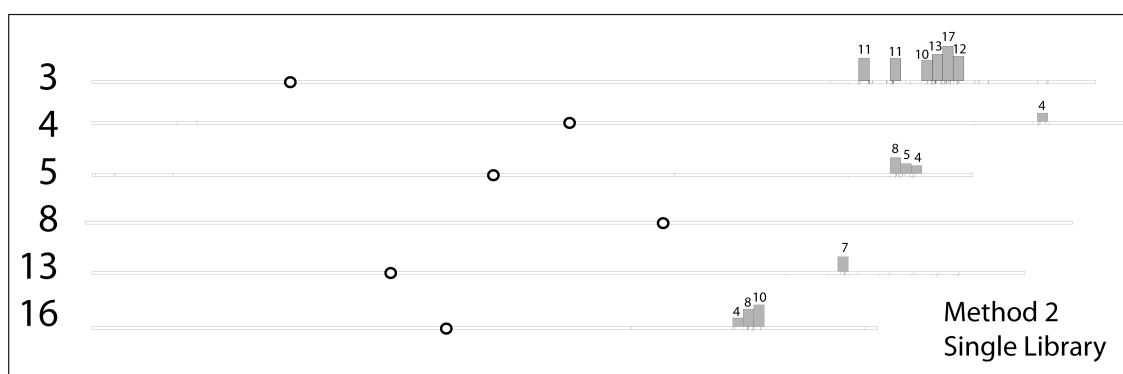


**Figure 1.** Chromosomal positions of Affymetrix SFPs and Goldengate SNPs identified between Clark and IsoClark. Chromosomes are labeled at the top according to number and centromere positions are shown as open circles. Red lines indicate the physical map positions of Affymetrix SFPs and blue lines indicate the physical map positions of Goldengate SNPs. Genomic regions co-incident for both SFPs and SNPs are indicated with yellow boxes and genomic regions exhibiting only Goldengate SNPs are indicated with clear boxes.

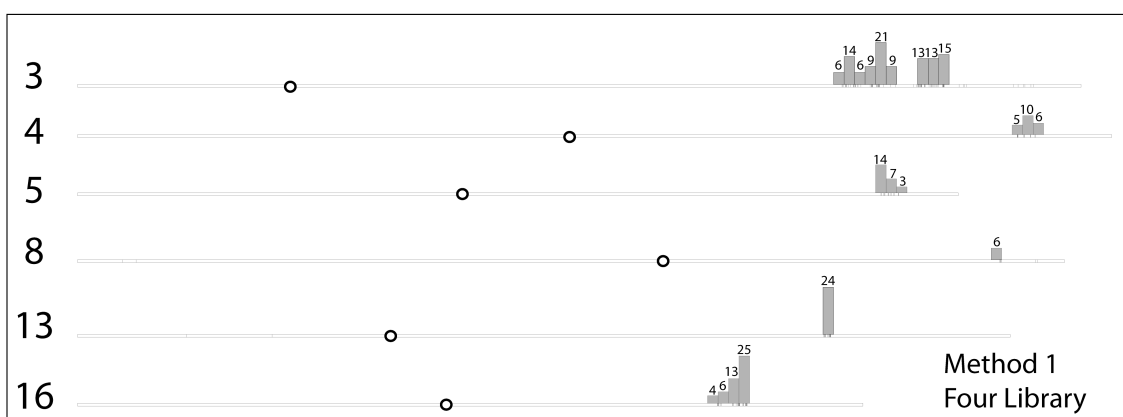
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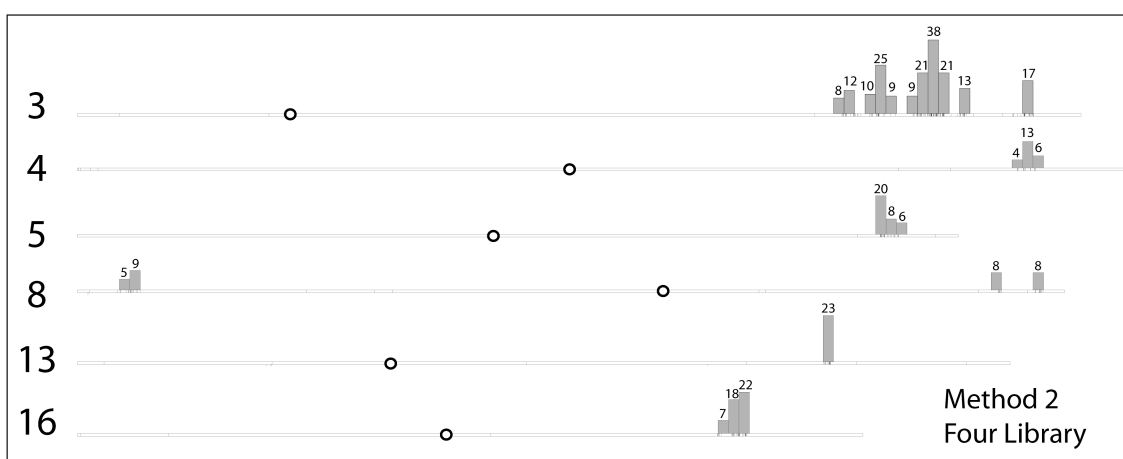
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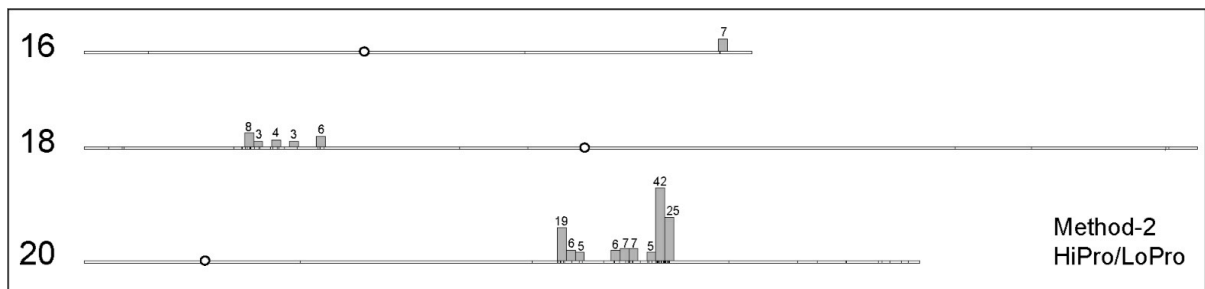
C



D



**Figure 2.** Significant intervals of SNP clustering between the Clark and IsoClark lines were found on six chromosomes: 3, 4, 5, 8, 13 and 16 as determined from the bootstrap method. Chromosomes are labeled on the left according to number and centromere positions are shown as open circles. Vertical boxes indicate 500,000-nucleotide intervals. The number of SNPs found in each interval is indicated above the interval. A) Clustering of SNPs obtained from the 10-day root data using Method 1 on the single-library comparison. B) Clustering of SNPs obtained from the 10-day root data using Method 2 on the single-library comparison. C) Clustering of SNPs obtained from the 19-day root and leaf data using Method 1 on the four-library comparison. D) Clustering of SNPs obtained from the 19-day root and leaf data using Method 2 on the four-library comparison.



**Figure 3.** Significant clusters of SNPs for the seed protein lines were found on three chromosomes: 16, 18 and 20 as determined from the bootstrap method. Chromosomes are labeled on the left according to number and centromere positions are shown as open circles. Vertical boxes indicate 500,000-nucleotide intervals. SNPs were identified via Method 2. The number of SNPs found in each interval is indicated above the interval. SNPs were clustered from the seed-protein data that contained 14 libraries for each NIL.



## Supplemental Material

\* Material located in Appendix pg.118

### Supplemental Table S1

SNPs identified by Method 1 on the single-library comparison.

### Supplemental Table S2

SNPs identified by Method 2 on the single-library comparison.

### Supplemental Table S3

SNPs identified by Method 1 on the four-library comparison.

### Supplemental Table S4

SNPs identified by Method 2 on the four-library comparison.

### Supplemental Table S5

SNPs identified by Method 2 on the seed protein NIL pair.

### Supplemental File S1

Script used to determine intervals of significant SNP density based on a bootstrap method.

## **Chapter 4. Identification of candidate genes underlying an iron-efficiency QTL in soybean**

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## Abstract

Prevalent on calcareous soils in the United States and abroad, iron deficiency is among the most common and severe nutritional stresses in plants. In soybean commercial plantings, identification and use of iron-efficient genotypes has proven to be the best form of managing this soil-related plant stress. Previous studies conducted in soybean identified a significant iron-efficiency QTL explaining > 70 % of the phenotypic variation for the trait. In this research, we identified candidate genes underlying these QTL through molecular breeding, mapping and transcriptome sequencing. Introgression mapping was performed using two related near-isogenic lines in which a region located on soybean chromosome 3 required for iron-efficiency was identified. The region corresponds to the previously reported iron-efficiency QTL. The location was further confirmed through QTL mapping conducted in this study. Transcriptome sequencing and qRT-PCR identified, two genes encoding transcription factors within the region that were significantly induced in soybean roots under iron stress. The two induced transcription factors were identified as homologs of the subgroup Ib bHLH genes in that are known to regulate the strategy I response in *A. thaliana*. Re-sequencing of these differentially expressed genes unveiled a significant deletion within a predicted dimerization domain. We hypothesize this deletion disrupts the FIT / bHLH heterodimer that has been shown to induce known iron acquisition genes.

## Introduction

Iron deficiency chlorosis (IDC) is a worldwide concern. In soybean a 20% yield reduction has been reported for each one-point increase in iron chlorosis score (Froehlich and Fehr, 1981). Hansen et al. (2003) estimated an annual loss of 120 million dollars due to IDC in the United States alone. Soybean bushels are selling approximately seven dollars higher today than in 2004, increasing the estimated current revenue losses to over 260 million dollars.

Iron is one of seven essential micronutrients required for plant growth (Crosa, 1989). If concentrations of micronutrients within the plant are too high, they become toxic and will stunt growth in the same fashion as an insufficiency (Havlin et al., 1999). Iron deficiency is one of the most common micronutrient deficiencies in the world due to alkaline soil conditions ( $\text{pH} > 8$ ) that maintain ferric ( $\text{Fe}^{3+}$ ) iron in an insoluble state unavailable to some plant genotypes. Conversely, most plants are able to uptake and utilize soluble ferrous ( $\text{Fe}^{2+}$ ) iron, prevalent in soils with pH ranging from 6.5 to 7.5 (Havlin et al., 1999). Iron deficiency in plants is commonly an issue of iron availability and not one of iron supply, as iron is the fourth most abundant element on the earth's surface.

Iron deficiency chlorosis (IDC) is caused by a decrease in photosynthetic pigments (Chlorophylls a/b, Carotene, Xanthophylls, etc.), in addition to a reduced electron

transport potential and a significant reduction in the thylakoid system (Spiller and Terry, 1980; Terry, 1980; Taylor et al., 1982). Symptoms of IDC vary from interveinal yellowing to complete necrosis (Cianzio, 1979; Havlin et al., 1999), which most genotypes overcome at the end of the growing season, however, severity of chlorosis is tightly correlated with end-of-season yield losses (Inskeep and Bloom, 1987).

The best way to manage IDC is through the use of Fe efficient varieties in commercial plantings. This is the reason why so much research has been devoted to identifying genetic markers associated with iron-efficiency for use in breeding programs. Work by Lin et al. (1997), showed that there are two genetic models to explain IDC resistance at the molecular level, which had been previously identified by classical genetic studies (Cianzio et al., 1980; Cianzio and Fehr, 1982). One model involves a single major gene with modifiers (Cianzio et al., 1980) identified in the Anoka x A7 population, the associated QTL was located on chromosome 3 explaining >70% of the phenotypic variation (Lin et al., 1997). The second model involved multiple genes (Cianzio and Fehr, 1982) identified in the PrideB216 x A15 population which represented a classic quantitative mode of inheritance. Lin et al., (1997) was unable to identify QTL in this population.

All plants except the Poaceae family have adapted what is known as the Strategy I response (Marschner et al., 1986) in order to reduce and acquire  $\text{Fe}^{+2}$  from soils. In

Strategy I, plants pump protons across the plasma membrane via an H<sup>+</sup>-ATPase to acidify the surrounding soil (Fox and Guerinot, 1998). The resultant acidification helps solubilize unavailable ferric iron. A ferric-chelate reductase (FRO2) reduces chelated Fe<sup>3+</sup> into available Fe<sup>2+</sup> (Walker and Connolly, 2008). The reduced iron (Fe<sup>2+</sup>) is transported across the plasma membrane via an iron-regulated transporter (IRT1) (Korshunova et al., 1999). Recently, work in *Arabidopsis thaliana* and *Lycopersicon esculentum* has made great strides at elucidating plant responses to iron limiting conditions. The response is initiated by an unknown factor that results from the recognition of low cellular iron, which then activates transcription factors AtFIT, AtbHLH038, and AtbHLH039 (Bauer et al., 2007; Wang et al., 2007). Yuan et al. (2008) showed that in *A. thaliana* AtFIT/AtbHLH038 or AtFIT/AtbHLH039 was capable of initiating GUS expression when regulated by AtFRO2 and AtIRT1 promoters. Their results suggested that AtFIT binds as a heterodimer to either AtbHLH038 or AtbHLH039 to initiate expression of AtFRO2 and AtIRT1 under iron limiting conditions (Yuan et al., 2008).

Soybeans have a production disadvantage when they are unable to evoke an effective iron deficiency stress response. Near-isogenic lines (NILs) differing in their response to iron were developed by the USDA-ARS in the early 1970's (Bernard, 1975) and have become some of the standard lines for IDC research in soybean. From this work, two lines identified as PI 547430 and PI 547698 resulted from the

cross between Clark (PI 548533)(6) x T203 (PI 54619) and Harosoy (PI 548573)(6) x T203 (PI 54619) respectively.

Introgression mapping is a viable method for delineating the genetic differences between NILs (Nichols et al., 2006; King et al., 2007). Nichols et al. (2006) showed that fine mapping of NILs could identify introgressed regions in chromosomes. They further showed that additional rounds of backcrossing could narrow the introgression and help identify candidate genes. More recently, SNPs generated from next generation sequencing of NILs have been used effectively to map introgressions (Severin, Peiffer et al., 2010). Severin and Peiffer et al. (2010) identified multiple genomic regions from T203 that were back-crossed (introgressed) into PI 547430 (Clark background). An introgression on chromosome 3 was significantly larger than the other small introgressions scattered throughout the genome. It is also noteworthy that the Gm03 introgression identified within the NILs using SNPs from next generation sequencing is positionally coincident with the position of the major QTL identified in the Anoka x A7 population (Lin et al., 1997; Severin, Peiffer et al., 2010). Mapping the T203 introgression in the other T203 derived NIL, PI 547698 (Harosoy background), will confirm the importance of this genomic location for iron-efficiency.

A microarray study in the Clark / PI 547430 NILs revealed that the number of induced genes in Clark under iron limiting conditions was more than four times that

of the iron-inefficient PI 547430 (O'Rourke et al., 2009). Furthermore, the induced genes cluster in the genome, however the clusters did not correlate with any of the 11 previously identified QTL (Lin et al., 1997; O'Rourke et al., 2009). For this reason it was hypothesized a gene(s) within the known iron-efficiency QTL acts transiently to initiate the iron stress response of genes residing outside of the known QTL region: presumably a transcription factor.

In this paper, we identify a region on chromosome 3 that is required for iron-efficiency and is shared between multiple populations. Candidate genes within the region as well as gene expression differences between near-isogenic lines were identified. Furthermore we characterize potentially significant alleles between iron-efficient and iron-inefficient lines. We also discuss how these alleles may alter gene function and induce an iron-inefficient response in an otherwise efficient background

## Results

### *Identification of the T203 introgressions in two NILs*

USDA-ARS scientists generated NILs by crossing the iron-efficient recurrent parents, Clark (PI 548533) and Harosoy (PI 548573) with an inefficient donor parent, T203 (PI 54619). The resulting F2 progeny were screened for iron-efficiency then backcrossed to the recurrent parent for five generations to create the iron-inefficient NIL. The genetic material from T203 introgressed into each efficient background



therefore presumably contains the gene(s) required to induce an iron-inefficient response in an otherwise efficient background (Bernard, 1975).

Identification of the introgressed DNA within the NIL genome was accomplished through a genome-wide survey using greater than 860 published simple sequence repeat markers (SSRs), available at Soybase.org (composite map 2003, Soybase.org). At least one marker representing the allele from the donor parent (T203) was detected on 12 of the 20 chromosomes for each background, Clark and Harosoy. Only on chromosome 3 (Gm03) was an introgressed region common for the two NILs (Table 1). PI 547430 and T203 shared alleles for markers Satt387 -> Sat\_125 and PI 547698 and T203 shared alleles for markers Sat\_275 -> Satt339 (Figure 1). In each NIL the region contained ~37% of all the introgressed markers.

A comparison of the overlapping region between the two pairs of NILs allowed for the region of interest to be narrowed to the 4.2 Mbp overlap (Satt387 -> Satt339) (Figure 1). Since, there were no other overlapping regions on any of the other chromosomes, it is likely that the gene(s) controlling iron-efficiency within the Clark/Harosoy NILs resides within this region. The region of introgression overlap on Gm03 correlates with the location of a known iron-efficiency QTL in an A7 x Anoka population (Lin et al., 1997; Lin et al., 2000). The iron-inefficient parent of this population, Anoka, is the result of a three-way cross between (Lincoln<sup>2</sup> x Richland) x Korean. It is important to indicate that Clark, one of the parents of the NIL, also

originated from the same single cross of (Lincoln<sup>2</sup> x Richland), later used as a parent in the tree-way cross from which Anoka was developed. Although Anoka was not generated in the same fashion as the two NILs, markers screened against Lincoln, Richland, Korean, and Anoka identified a region corresponding to the NIL introgression in which Anoka and Korean shared marker alleles (data not shown), suggesting transfer of genetic material within this region from the inefficient parent, Korean, to Anoka.

#### *High density mapping of the Anoka x A7 QTL*

Integrating the genetic map developed by Lin et al. (1997) with previously mapped SSRs and newly available BARCSOYSSRs (Song et al., 2010) resulted in a higher density genetic map. The 1993 and 1994 combined visual phenotypic data collected by Lin et al. (1997) was reanalyzed using the integrated map. The QTL on Gm03 was reconfirmed with a LOD score of 15.29, which is significantly higher than the permuted genomewide LOD cutoff value of 3.8 (Figure 2). The QTL region identified in 1997 spanned Gm03 from Sat\_33 -> BLT\_15. By placing additional markers on the map the predicted QTL region on Gm03 was significantly narrowed using the integrated map (Sat\_33 -> S03\_1129) (Figure 2).

#### *Reducing the donor parent introgression in PI 547430*

Following the identification of the T203 introgressed region in the iron-inefficient NIL PI 547430, further backcrossing to the recurrent parent (Clark) resulted in a BC6

population of PI 547430 during the summer of 2009 (Ames, IA). Two hundred cross pollinations were performed and 156 resulted in seed set. Seed from the crosses were planted in the greenhouse during the winter 2009/2010. Four SSR markers within the introgressed region were used for screening, to distinguish between plants that resulted from crosses or self-pollinated seeds. From these checks, 98 F1 plant/F2 seed families were confirmed as true hybrids. In summer 2010, 40 seed of each confirmed plant/family were planted at Bruner farm (Ames, IA).

Leaf tissue was harvested early in the growing season, at plant stage V2, in order to genotype the nearly 6000 plants in the field. An initial round of genotyping was performed with four markers to identify plants with crossovers within the introgressed region (Sub-NILs). From this initial screen, 480 plants were identified with a recombination within the region and chosen for additional marker analysis to identify potentially narrowed introgression regions. Twenty-nine plants were identified as fitting into 10 different sub-NIL classes, segmenting the introgressed region (Figure 3).

#### *Phenotypic analysis of lines with narrowed introgressions*

Phenotyping of the lines representing various recombinant classes was performed during winter 2010/2011. Plants were scored for iron chlorosis symptoms using the standard visual 1-5 scale and a Minolta SPAD-502 chlorophyll meter. Two replications, with eight plants each, were grown hydroponically in iron limiting

conditions in the greenhouse. The recombinant classes segregated into two groups, those with introgressions originating from the left and those originating from the right side of the original PI 547430 introgression (Figure 3). Visual and SPAD score averages for each of the two groups and controls (Clark and PI 547430) were compared. Interestingly, the visual score and SPAD averages for all categories containing the donor parent allele for Satt387 were not significantly different from the averages for PI 547430 (P-value: 0.7366 / 0.8202) allele. Visual and SPAD averages for all classes containing the Clark allele for Satt387 were not significantly different from the averages of Clark (P-value: 0.0864 / 0.3051). Of the classes containing the donor parent allele for Satt387 class 1 contains the least amount of T203 derived DNA. Recombinant class 1 was not phenotypically different from the other introgressions containing the T203 allele for Satt387 (p-value = 0.097). Thus, we could narrow the donor parent introgression in PI 547430 from ~ 6 million down to just over 1 million base pairs.

#### *Fine mapping recombinant class 1*

Recombinant class 1, consisting of the F<sub>2:3</sub> family 4-25, contained the least amount of T203 alleles conferring iron-inefficiency. Eight plants from the 4-25 family were genotyped with 17 additional SSR markers to identify more precise recombination intervals. Five shared a recombination interval of S03-1141 -> S03-1155 (Figure 4). Three shared an even narrower recombination interval between S03-1110 and S03-1112 (Figure 4). The average phenotypic score for the two subgroups of plants

within class 1 were not significantly different from each other ( $p$ -value = 0.2552).

The introgression was therefore further narrowed to the ~250kb region identified by the three 4-25 plants with the recombination interval of S03:1110 -> S03:1112. The narrowing of this introgression significantly reduced the number of candidate genes from hundreds of genes down to eighteen genes potentially controlling the iron stress response in the NILs.

#### *Next-generation sequencing, RNA-Seq, analysis of NILs*

RNAseq was performed in replicate on RNA isolated from Clark leaves and roots 24 hours post iron stress. Eight libraries were constructed and sequenced producing 165,968,390 reads. The reads were then mapped to the soybean gene calls using Bowtie / Tophat (Trapnell et al., 2009). Differential gene expression between sufficient and iron insufficient conditions was determined using an FDR threshold of 0.05 in Cuffdiff (Roberts et al., 2011). The eighteen genes within the narrowed Gm03 introgression were examined in the RNA-seq data to determine their expression differences between iron sufficient and iron insufficient conditions (Table 2). In both replicates only two of the 18 genes were significantly differentially expressed in the roots, Glyma03g28610 and Glyma03g28630. The two genes share 87.8% peptide similarity and are homologous to the *Arabidopsis thaliana* AtbHLH038 with 66.7 and 63.9% peptide similarity respectively. The expression of the genes was significantly increased ( $p < 0.05$ ) under iron stress conditions compared to that of the iron-sufficient condition. Expression of the two genes was also restricted to the

roots and expression was not detected in leaves of plants grown in stressed or non-stressed conditions. The leaves showed differential expression of six genes within the identified introgression (Glyma03g28450, Glyma03g28490, Glyma03g28590, Glyma03g28480, Glyma03g28510 and Glyma03g28570) in one of the two biological replicates but not in the other. When the two replicates were combined, four of the genes were confirmed as differentially expressed (Glyma03g28450, Glyma03g28490, Glyma03g28480 and Glyma03g28570) (Table 2). Within the narrowed introgression four genes were identified as transcription factors (Glyma03g28610, Glyma03g28630, Glyma03g28500, and Glyma03g28440) (Wang et al., 2010).

#### *Gene expression comparison between Clark and PI 547430*

Previous studies in *A. thaliana* have shown that AtbHLH038 is induced under iron limiting conditions (Yuan et al., 2008). In our study, homologs of this gene (Glyma03g28610/28630) were also induced. qRT-PCR was used to confirm the expression pattern of these genes identified through RNA-seq analysis. qRT-PCR confirmed the results of the RNA-seq. Both genes showed a greater than two-fold induction under Fe-insufficient conditions (Table 3). While RNA-seq data was not generated for PI 547430, qRT-PCR expression analysis was performed on the same genes in the inefficient NIL. A greater than two-fold induction of Glyma03g28610 and Glyma03g28630 was observed in PI 547430 under iron stress conditions (Table

3). Clark and PI 547430 did not show a significant difference in the expression levels of these genes under iron stress conditions.

In *A. thaliana*, it has been shown that AtFIT binds AtbHLH038 as a heterodimer to activate AtFRO2 and AtIRT1 (Yuan et al., 2008). We used the available RNA from Clark and PI 547430 to investigate the expression patterns of these downstream genes. We identified the soybean homologs of the *A. thaliana* genes using peptide similarity and found that Glyma07g07380 (GmFRO2) is homologous to AtFRO2 with 72.5% peptide similarity. Glyma07g34930 (GmIRT1) is homologous to AtIRT1 with 77.7% peptide similarity, and Glyma12g30240 (GmFIT) is homologous to FIT with 62.1% peptide similarity. Using primers designed from these genes qRT-PCR showed that Clark induced Glyma03g28610, Glyma03g28630, GmFIT, and the two downstream genes GmFRO2 and GmIRT1 under iron stress (Table 3). PI 547430 had significant induction of GmIRT1 yet failed to induce GmFIT or GmFRO2 under iron stress even with high Glyma03g28610 and Glyma03g28630 expression (Table 3). The expression levels of GmFIT, GmFRO2 and GmIRT1 in Fe stressed Clark were significantly greater (3, 5 and 2.5-fold respectively) than that of PI 547430 under Fe stress.

#### *gDNA sequencing of differentially expressed transcription factors*

Glyma03g28610 and Glyma03g26830 were the only two transcription factors found to be differentially expressed within this region. These two genes were sequenced

in Clark, Anoka and T203 to determine if allelic differences could be identified through sequencing analysis. Allelic differences were identified between the iron-efficient Clark and the iron-inefficient T203 and Anoka. A twelve base pair deletion was identified in Glyma03g28610 (Gm03:36552946..36552957) an in-frame four amino acid deletion (PELQ). Primers were designed to identify the presence or absence of the deletion and used to screen Harosoy, Lincoln, Richland, A7, Korean, PI 547698 and PI 547430. The twelve base pair deletion was identified only in the iron-inefficient cultivars (Anoka, Korean, T203, PI 547698 and PI 547430). It was not detected in the iron-efficient genotypes (Clark, Harosoy, Lincoln, Richland and A7) (Figure 5).

## Discussion

### *Whittling donor parent introgressions*

In this report we have narrowed a large list of candidate genes derived from a classical genetic QTL study down to a manageable list of eighteen genes. Two of the candidate genes are homologous to genes in *A. thaliana* known to play a major role in iron acquisition. We used a variety of mapping techniques and additional backcrossing of NILs to narrow the region of interest on chromosome 3 down to a 250 kb region. The research reported herein shows that using the whole genome sequence of soybean, classical genetics and genomic technologies, we were able to



identify the genes underlying a QTL previously identified by Lin et al. (1997), nearly 15 years ago.

We investigated two sets of near isogenic lines sharing a common donor parent and identified regions of the genome introgressed from the donor parent into each of the recurrent parents. The results reported here using SSR markers to delineate the T203 introgressions in the PI 547430 NIL confirmed the introgressions using SNPs reported by Severin et al. (2010). However, by also mapping T203's introgression into Harosoy we were able to significantly decrease the proportion of the introgressed region likely to contain the gene(s) involved in the iron stress response. The use of the common donor parent allowed for the overlay of the two maps. Fortunately, the introgressed regions in each NIL shared only a small overlap, which allowed for further dissection of this region down to 4.2 Mb (Satt387 -> Satt339).

Early IDC QTL studies used the mapping population Anoka x A7 to identify an iron-efficiency QTL on chromosome 3 that was responsible for > 70% of the phenotypic variation (Lin et al., 1997). Since the late 1990's the marker density on the soybean genetic map has been greatly improved. It was for this reason that the population used in this early QTL study was rescreened with the approximately 900 newly available SSR markers (Soybase.org). Using the original phenotypic data the major QTL on chromosome 3 was reconfirmed in our study and further refined. Lin et al. (1997) hypothesized that the QTL on chromosome 3 was the result of a single major

gene within the QTL responsible for their phenotypic observations. Our findings with the NIL mapping in both the Harosoy and Clark backgrounds support their hypothesis in addition to reconfirm the single major QTL in the Anoka x A7 mapping.

To identify the gene(s) responsible for Fe inefficiency within the QTL, it was necessary to narrow the list of candidate genes. Previous studies have shown that additional backcrosses in NIL populations allows for narrowing the introgression boundaries (Nichols et al., 2006). The successes seen in this approach led us to develop the sixth backcross of PI 547430 to the recurrent parent Clark during summer 2009. We identified two distinct phenotypic classes; those containing the PI 547430 allele for Satt387 that displayed IDC symptoms similar to that of PI 547430 and those with the Clark allele for Satt387 that were healthy. We were further able to delineate the region with the use of 17 previously unmapped BARCSOYSSRs (Song et al., 2010). Recombinant class 1 harbored plants with two distinct recombination intervals not significantly different from each other with respect to IDC response. From this we were able to narrow the genetic interval further to a manageable 250 kb region significantly reducing the number of candidate genes from > 800 down to a mere 18. The 250kb region co-locates with the Anoka x A7 QTL to the centromeric side of the introgression overlap indicating that the Anoka x A7 and the two NILs may share common alleles for a gene(s) facilitating the IDC response.

### *Candidate gene expression*

O'Rourke et al. (2009) performed microarray studies using the Clark/PI 547430 NIL and hypothesized that regulatory elements within the known iron QTL were responsible for gene expression changes of known iron genes located outside the QTL region. Our study supports that hypothesis. Four of the eighteen genes identified within the narrowed introgression were annotated as transcription factors Glyma03g28440, Glyma03g28500, Glyma03g28610, and Glyma03g28630 (Wang et al., 2010). Two of these genes Glyma03g28610 and Glyma03g28630 are interesting as they are homologs of *A. thaliana* AtbHLH038. AtbHLH038 is a member of a subgroup of bHLH genes known as lb bHLH genes and have been shown to be induced upon iron stress (Yuan et al., 2008). It has been proposed that AtbHLH038 can bind as a heterodimer to AtFIT to enhance the expression of the downstream iron acquisition genes, AtFRO2 and AtIRT1 (Yuan et al., 2008). Polymorphisms in these two transcription factors likely underlie the QTL in question. Of the 18 genes identified within the narrowed region of interest, Glyma03g28610 and Glyma03g28630 were induced upon iron stress and were the only two genes found to be significantly differentially expressed in the roots and the only two transcription factors differentially expressed in either the roots or the leaves. qRT-PCR confirmed the expression trends of Glyma03g28610 and Glyma03g28630 in addition to the soybean homologs of AtFIT, AtFRO2 and AtIRT1. LeFER, the AtFIT homolog in *L. esculentum* is not induced upon iron stress (Ling et al., 2002) whereas AtFIT is induced upon iron stress along with AtbHLH038/039, which initiate expression of

AtFRO2 and AtIRT (Yuan et al., 2008). We have shown that FIT expression in soybean corresponds with the induction observed in *A. thaliana* and not *L. esculentum* as soybean induces GmFIT expression upon iron stress.

Transcriptome analysis was not conducted on PI 54730, however by using qRT-PCR we were able to investigate the inefficient NIL's response to iron stress at the gene expression level. PI 547430 showed no significant change in GmFIT or GmFRO2 expression between iron sufficient and iron insufficient conditions, however GmIRT1 was still induced greater than two fold though significantly less than that of Clark. AtIRT1 is typically thought to have coordinate expression with AtFRO2 (Connolly et al., 2003), however our results showed a significant difference in the coordinated induction of GmFRO2 and GmIRT in PI 547430 under iron limiting conditions. The lack of GmFRO2 induction is quite significant since the reduction of  $\text{Fe}^{3+}$  to  $\text{Fe}^{2+}$  at the rhizosphere has been shown to be a rate-limiting step for iron uptake (Grusak et al., 1990; Connolly et al., 2003).

Although not analyzed in depth in this study, six genes were differentially expressed in Clark leaves. Of these genes Glyma03g28570 stands out. It is annotated in the KEGG database (Kanehisa and Goto, 2000) as a member of the two-component response regulator ARR-A family. To et al., (2004) identified the Type-A *Arabidopsis* response regulators as having a negative role in the regulation of cytokinins. Recently, Seguela et al. (2008) identified for the first time that cytokinins were

involved in regulation of AtFIT, AtFRO2 and AtIRT1. Potentially Glyma03g28570 is another one of the underlying genes behind the significant effects of this iron-efficiency QTL.

Interestingly, in the Anoka x A7 mapping population the iron-inefficient parent Anoka was derived from a cross between (Lincoln<sup>2</sup> x Richland) x Korean. Clark, the iron-efficient recurrent parent used in the earlier NIL mapping, was also derived from a Lincoln<sup>2</sup> x Richland cross. From this we would hypothesize that Korean donated the iron-inefficient allele(s) in Anoka. We mapped two distinct blocks of SSRs containing alleles derived from Korean on chromosome 3 which corresponded with the overlapped introgression identified by both of the NILs (data not shown). The identification of the same genomic region in the NIL and QTL mapping studies along with the twelve base pair deletion in Anoka and Korean strongly indicates Glyma03g28610 or another gene(s) within this region as having a major effect on the iron-efficiency of soybean. Furthermore, if the dimerization capabilities of this AtbHLH038 homolog are in fact hindered by the twelve base pair deletion it could explain why plants harboring this mutation aren't as effective at surviving in low iron environments.

#### *Mutant identification*

The lack of a significant gene expression change for Glyma03g28610 and Glyma03g28630 between Clark and PI 547430 under iron stress conditions indicates

that if these genes are the underlying cause of the QTL then the difference between the lines may be due to gene function. We identified two alleles of Glyma03g28610 that had 100% correlation with iron-efficiency in this study. The iron-inefficient cultivars Anoka, T203, PI 547430, PI 547698, and PI 548360 all shared a twelve base pair deletion within the second exon of this gene. The deletion is an in-frame 4 amino acid deletion that doesn't appear to hinder accumulation of the gene's transcript. According to NCBI's Conserved Domain Database the deletion spans one of the 14 predicted dimerization sites and shifts two others (Figure 6). This deletion could potentially alter Glyma03g28610's ability to bind as a heterodimer with GmFIT, or alter the confirmation of the protein so that it is no longer able to bind the e-box within the promoter regions of GmFRO2 and GmIRT1. This could explain why GmFRO2 is not induced in PI 547430 upon iron stress. It has been hypothesized that there is an additional regulatory step between AtFIT and AtIRT1 (Colangelo and Gueriot, 2004). This unknown factor could be the cause of the GmIRT1 expression in PI 547430. In addition, it has been shown that AtFIT regulates AtFRO2 at the level of transcription and IRT1 at the level of protein accumulation (Colangelo and Gueriot, 2004). It was beyond the scope of this study to look at protein accumulation. Thus, even though we observed an induction of GmIRT1 mRNA we cannot claim that there is an increase of GmIRT1 protein in the soybean roots. PI 547430 lacked the ability to induce GmFIT under iron limiting conditions, which indicates that there may be further unknown factors within the iron response pathway capable of inducing GmIRT1.

## Conclusion

In this report, we show that mapping sites of introgressions in multiple NILs sharing a common donor parent is a viable method for identification of genomic regions controlling phenotype. Furthermore, through the creation of Sub-NILs and gene expression analysis that we were able to identify two candidate genes within the NIL introgression on chromosome 3 that co-localizes with a known Fe-efficient QTL in the Anoka x A7 population. Finally, we identified a 12 bp deletion in one of the candidate genes that is shared in the inefficient lines in both NILs and the inefficient parent, Anoka, in the Anoka x A7 population.

## Materials and Methods

### *Introgression mapping*

Lines used in this study were derived from backcrossing the two efficient recurrent parents Clark and Harosoy crossed with the common donor parent T203 (Bernard, 1975). Mapping of the T203 (donor) introgression into both PI 547430 (Clark background) and PI 547698 (Harosoy background) utilized 869 and 864 SSR markers respectively. Markers were used to amplify DNA from the donor, and recurrent parents, and from the isolines. PCR product was visualized on a 6% polyacrylamide gel. Screened markers were categorized into three classes: not informative, informative between the parents, and introgressed DNA. The

introgressed class was defined by the isolate's banding pattern matching that of the donor parent. All markers had a known location on the whole genome sequence (Glyma1.01) ([www.SoyBase.org](http://www.SoyBase.org)) and thus identified regions of introgression into the near isogenic line.

#### *QTL mapping in the Anoka x A7 population*

Plant material for QTL mapping originated from 92  $F_{2:4}$  lines from the Anoka x A7 population, which was previously used to map iron deficiency chlorosis QTL (Lin et al., 1997). The genetic linkage map was created by screening the population with 916 known SSR markers ([SoyBase.org](http://SoyBase.org)) and an additional 303 untested BARCSOYSSR markers (Song et al., 2010). The marker scores for each line were then imported into Mapmaker 3.0 (Lander et al., 1987). A logarithm of the odds (LOD) threshold of 4.0 was used in a 3-point linkage analysis to construct the map.

Phenotypic data obtained by Lin et al. (1997) was used in the research reported here. Linkage maps were imported into MAPQTL6 (Van Ooijen, 2009), and QTL were mapped using interval mapping and multiple-QTL mapping (Lander and Botstein, 1989; Jansen, 1993, 1994). A significant experimentwise LOD threshold of 3.8 was determined by 1,000 permutations of the data.



### *Backcross advancement of PI 547430*

To further increase the chances of narrowing the introgressed region within the NIL (Clark background) an additional backcross generation was performed in these lines, during the 2009 summer. Seventy-five Clark and 75 PI 547430 seed were planted on May 19<sup>th</sup>, 2009 at the Bruner farm (Ames, IA). Two hundred crosses were performed which produced 156 seeds that were planted during the 2009 / 2010 winter season in the greenhouse at Ames, IA. The BC6 plants were screened with four SSR markers to confirm heterozygosity within the introgressed region. Forty seed from each confirmed cross were then planted at the Bruner farm (Ames, IA) on May 27<sup>th</sup>, 2010. Approximately 2600+ plants were screened with Satt387, Satt521, Sat\_091 and Sat\_295 to identify recombinations within the introgressed region. Of these, 480 plants with recombination within the introgression region were selected for genotyping with four additional SSR markers (Sat\_236, Satt549, Satt339, Sat\_304) to better delineate the recombination intervals.

### *Hydroponic phenotyping*

In the winter of 2010 seed of each line was sowed on germination paper for 7 days before seedlings transfer to a custom plant holder suspended over a 10L foil wrapped plastic bucket. Two replicates containing eight plants in each recombinant class were grown under iron stress conditions (50  $\mu$ M Fe(NO<sub>3</sub>)<sub>3</sub>•9H<sub>2</sub>O), following the procedure outlined by O'Rourke et al. (2007). Each 10L bucket contained: 2mM MgSO<sub>4</sub>•7H<sub>2</sub>O, 2mM Mg(NO<sub>3</sub>)•6H<sub>2</sub>O, 2.5mM KNO<sub>3</sub>, 1mM CaCl<sub>2</sub>•2H<sub>2</sub>O, 4mM

Ca(NO<sub>3</sub>)<sub>2</sub>•4H<sub>2</sub>O, 0.02mM KH<sub>2</sub>PO<sub>4</sub>, 542.5μM KOH, 217μM KOH, 20μM MnCL<sub>2</sub>•2H<sub>2</sub>O, 50μM ZnSO<sub>4</sub>•7H<sub>2</sub>O, 50μM CuSO<sub>4</sub>•5H<sub>2</sub>O, 0.2μM Na<sub>2</sub>MoO<sub>4</sub>•2H<sub>2</sub>O, 1μM CoSO<sub>4</sub>•7H<sub>2</sub>O, 1μM NiSO<sub>4</sub>•6H<sub>2</sub>O and 10μM H<sub>3</sub>BO<sub>3</sub> in accordance with research by Chaney and Bell (Unpublished Advisory Letter, 1989) and O'Rourke et al. (2007). A 3% CO<sub>2</sub>-air mixture was used to maintain a pH level of 7.8. Each day a stock solution containing 30M K<sub>2</sub>HPO<sub>4</sub>, 222M KH<sub>4</sub>NO<sub>3</sub>, and 0.179M H<sub>3</sub>BO<sub>3</sub> was added into the nutrient solution system to replace nutrients utilized by the growing plants. The greenhouse photoperiod was held constant at 14-hours with a 10-hour dark period for the entirety of the experiment. Plants were phenotyped 15 days after transfer into hydroponics at the V3 stage. The 1-5 visual IDC score identification was performed as outlined by Ciazio et al. (1979). In addition to visual score, SPAD readings were also recorded with a Minolta SPAD-502 chlorophyll meter (Tokyo, Japan), immediately following visual score determination.

*RNA sampling of Clark and PI 547430 tissues following iron-shock (root and leaf)*

Plants of Clark and PI 547430 were grown using a similar greenhouse hydroponic system with sufficient amounts of iron for 14 days according to the methods described above. After this period plant roots were rinsed six times to remove iron and other nutrients traces and transferred to either Fe-sufficient (100 μM Fe(NO<sub>3</sub>)<sub>3</sub>•9H<sub>2</sub>O) or Fe-insufficient (50 μM Fe(NO<sub>3</sub>)<sub>3</sub>•9H<sub>2</sub>O) hydroponic conditions. After 24 hours of being transferred, foliar and root tissue was collected and flash-frozen for storage. RNA was later extracted and purified using RNeasy mini kit

(QIAGEN Inc., Valencia, CA). The isolated RNA was then treated with DNase to remove final traces of DNA and stored at -80C until sequenced.

#### *Illumina Next-generation sequencing of RNA*

Four libraries were sequenced using a Genome Analyzer II (Illumina). The libraries consisted of two Fe-sufficient and two Fe-insufficient biological replicates each containing three pooled replicates. Sequencing was performed using established methods at the National Center for Genome Resources (NCGR). Severin et al. (2010) have outlined the techniques of RNA-Seq sequencing. In brief, RNA with a poly-A tail was isolated and size selected for ~500 bp fragments. PCR amplified products of the size-selected library were checked for quality and loaded onto an Illumina flow cell for 36 cycles.

#### *Analysis of Next-generation sequencing Data*

The 36 bp reads were aligned to the soybean genome transcript map (Gmax\_109, (Schmutz et al., 2010) using Bowtie/Tophat (Trapnell et al., 2009). Alignment was limited to a seed length of 28 bases with a mismatch limit of two. The alignments were further processed using Cufflinks/Cuffdiff v1.0.2 (Roberts et al., 2011).

Transcript abundances were determined in Cufflinks using the fragments per kilobase of exon per million fragments mapped (FPKM), similar to the method described in Mortazavi et al., (2008). Normalization of samples was performed using upper quantile normalization in order to better achieve estimates of lowly expressed

genes (Bullard et al., 2010). Cuffdiff was used to identify gene expression differences between sufficient and insufficient iron conditions (Roberts et al., 2011). Significant differential gene expression was determined using a false discovery rate of 0.05 with a minimum of 30 alignments at a locus for a valid significance testing and a significant p-value cutoff of 0.05.

#### *Gene expression analysis through qRT-PCR*

Primers were designed from protein coding sequences (CDS, Phytozome.net) using Primer3 as a primer design tool (Rozen and Skaletsky, 2000). Primers were designed to amplify ~100 bp fragments ideally spanning the intron/exon border to differentiate RNA and DNA amplified product. RNA amplification was performed using SuperScript III Platinum SYBR Green One-Step qRT-PCR amplification kit (Invitrogen). Reactions were performed on a Stratagene Mx3000p Real-Time PCR machine (Agilent Technologies, Inc, Santa Clara, CA). Reactions were run for 44 cycles at a temperature of 60°C. A standard curve was created using 2.5, 12.5, 25, 100 and 150ng RNA concentrations along with a no template control, and a no reverse transcriptase well for each primer pair.

ELF-1B was chosen as the reference gene for this study based on previous reports by Wang et al., (2009). For each plate one test primer and one reference gene primer were screened with two technical replicates for each of the three biological replicates. This design allowed for the comparison between the quantity values

(Rieu and Powers, 2009). qRT-PCR reactions were used for analysis only if the dissociation curve showed one peak and the no-RT and no-template reactions were greater than 5 cycles away from the other data points. The standard curve was accepted if the  $r^2$  value was greater than 0.985.

The relative quantities were determined by normalizing the values for the test primers to that of the reference gene (ELF-1B). The technical and biological replicates were then averaged and log base 2 transformed.

### *gDNA sequencing*

DNA was extracted from Clark (PI 548533), Harosoy (PI 548873), T203 (PI 54619), PI 547430, PI 547698, Anoka (PI 548508) and A7 (PI 596526) using a 96-well c-tab method outlined by Dietrich et al. (Dietrich et al., 2002). Primers were designed across the predicted gene calls starting ~500 bp upstream of the transcription start site and continuing ~500 bp past the transcription stop. Individual primers were designed to amplify ~750 bp fragments. The sequencing methods used are outlined in Severin, Peiffer et al. (2010). Cleanup of the PCR reaction utilized exonuclease 1 and shrimp alkaline phosphatase. Sequencing was performed using the BigDye V3.1 sequencing kit on a 50 cm 96-capillary Applied Biosystems 3730XL DNA analyzer (Applied Biosystems). The sequences from all eight genotypes were then aligned using Sequencher V4.9 (Gene Codes Corporation). The alignments were used to identify SNPs between the sequenced lines.

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## Author Contributions

Gregory A. Peiffer wrote the manuscript, conceived the hypothesis and experimental design through the introgression identification, narrowing and gene expression changes. Keith E. King and Nicholas C. Lauter performed Fe efficiency QTL analysis in the Anoka x A7 population. Andrew J. Severin contributed to the gene expression analysis. Shun Fu Lin generated the Anoka x A7 population and phenotypic scores. Silvia R. Ciazio and Randy C. Shoemaker provided guidance and editorial assistance for the manuscript.

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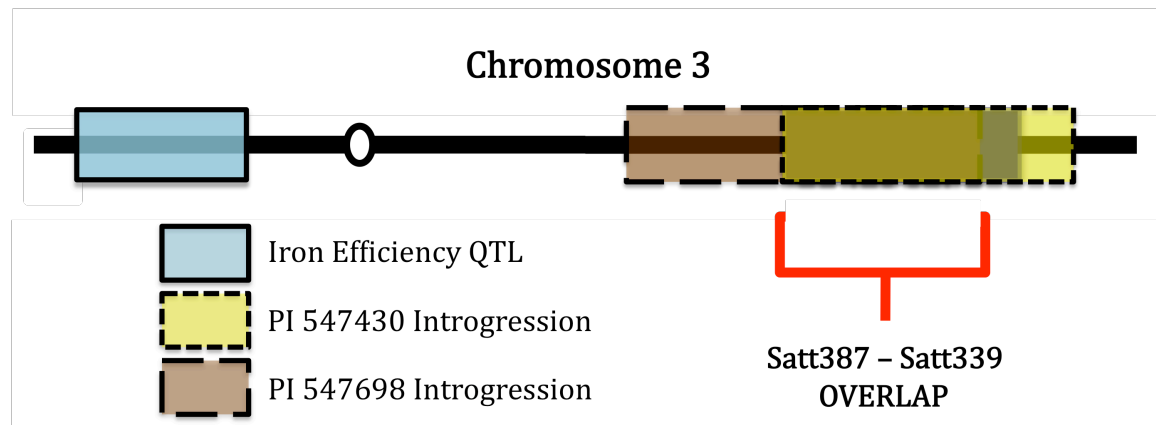
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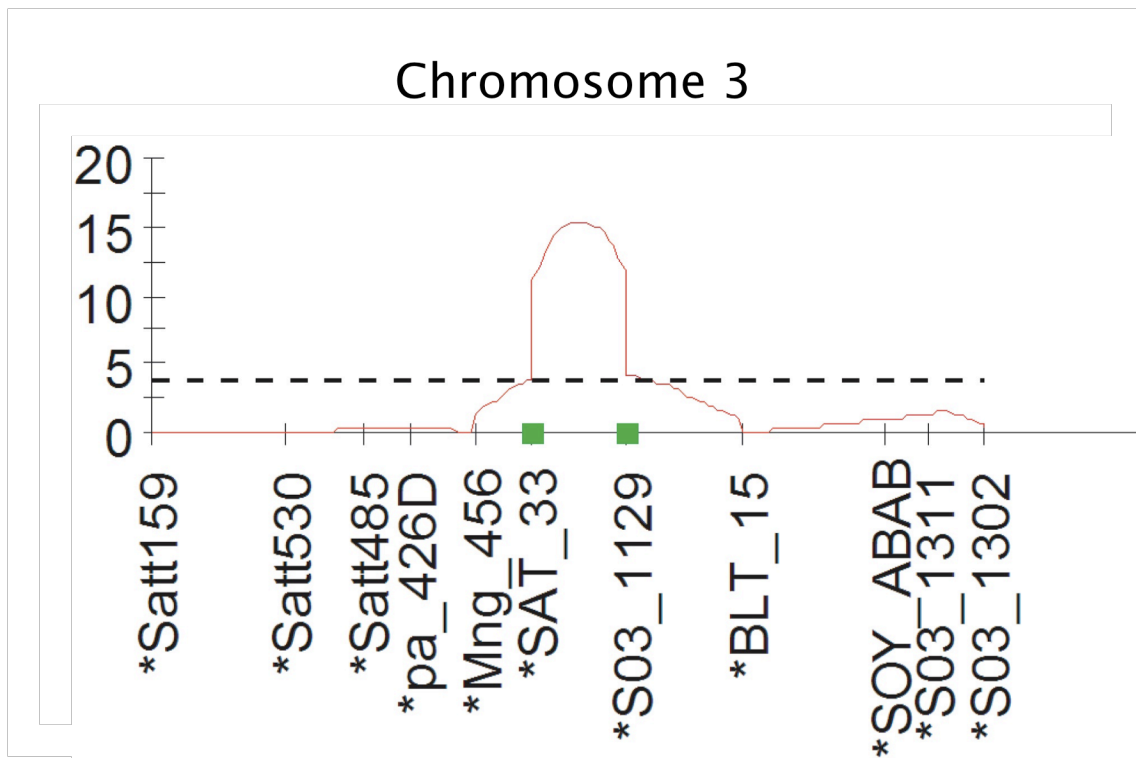
**Table 1.** Polymorphic SSR Markers Screened in the Two NILs

	Total Markers	Polymorphic Markers <sup>1</sup>	Introgressed Markers <sup>2</sup>	Percent on Gm03
Clark	869	415	27	37.7
Harosoy	864	345	19	36.9

Comparison of polymorphic markers between the two recurrent parents and T203. In each NIL approximately 37% of the polymorphic markers were clustered on chromosome 3. <sup>1</sup>Isoline same banding as Clark, <sup>2</sup>Isoline same banding as T203

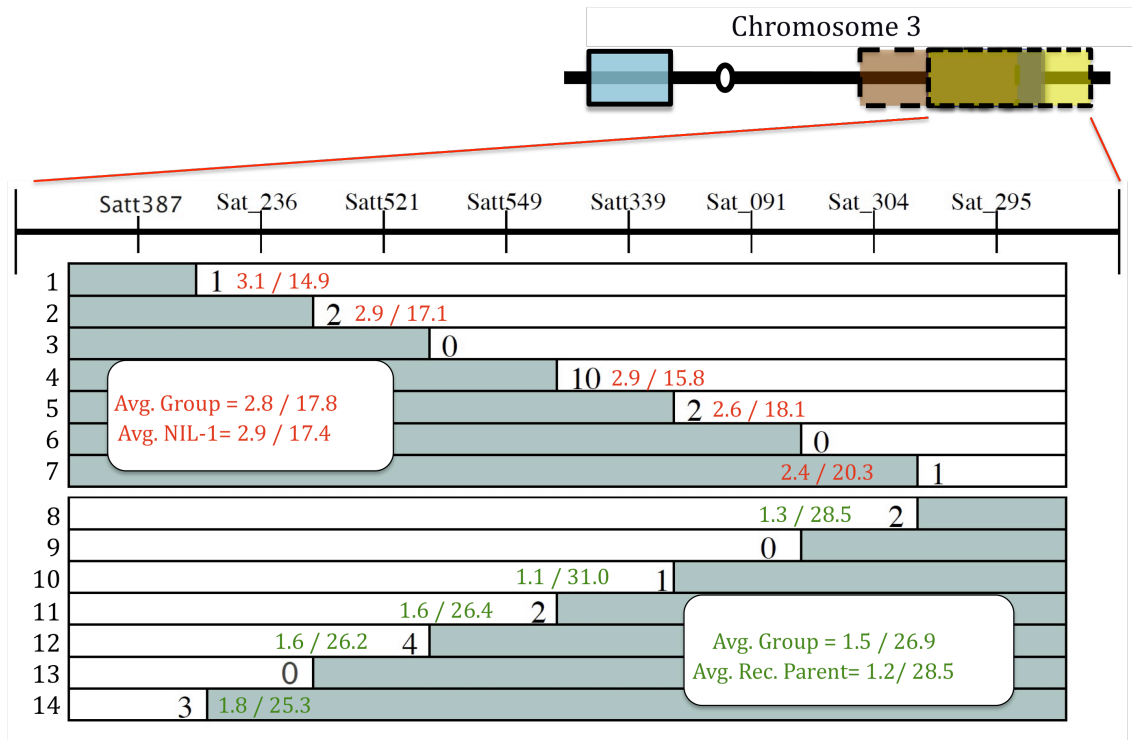


**Figure 1.** Overlay of the Two Near Isogenic Line introgressions and the iron efficiency QTL on chromosome 3. Blue solid boxes are the previously identified iron-efficiency QTL, the brown wide dashed box is the PI 547698 introgression, and the yellow short dashed box is the introgression in PI 547430.

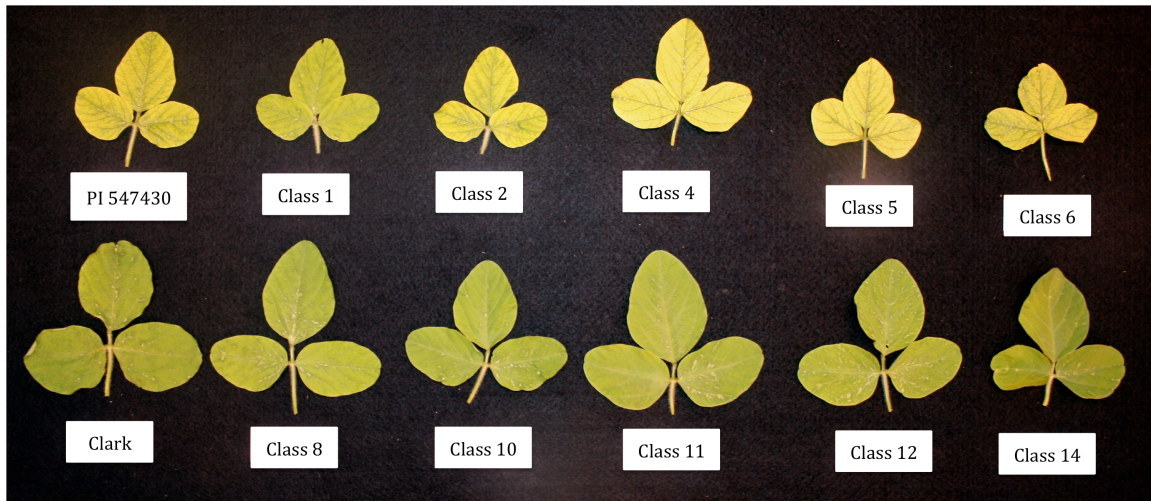


**Figure 2.** IDC visual scores mapped in the Anoka x A7 population for the combined 1993 and 1994 data. The only significant QTL was found on chromosome 3 with a LOD score of 15.29.

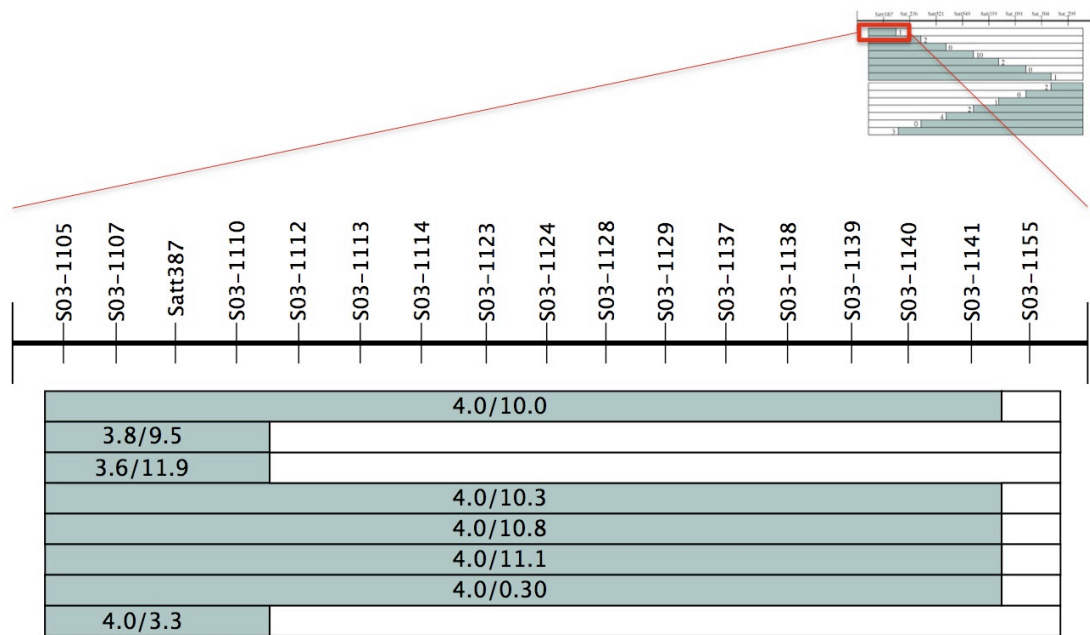
A



B



**Figure 3.** Sub-NIL phenotyping for in the PI 547430. A, Diagram of the Sub-NILs identified across the BC6 PI547430 introgression site. Phenotypes for each class are reported with the visual score / SPAD reading. Shaded region indicates donor parent alleles. B, Representative images of the plants in each class. Controls for each group are PI 547430 and Clark respectively.



**Figure 4.** Fine mapping of recombinant class 1. Shaded regions indicate donor parent alleles, the non-shaded regions indicate Clark alleles. The phenotypic scores are recorded with the visual score / SPAD score.

**Table 2.** Annotation of Genes within Narrowed Introgression  
 PFAM Annotations of the candidate genes within the whittled introgression on chromosome 3. Differential expression is indicated for each of the data sets.

<b>Gene Name</b>	<b>Differential expression</b>	<b>PFAM ID</b>	<b>PFAM Description</b>
Glyma03g28440		PF00651	BTB/POZ domain
Glyma03g28450	Leaf 1 & 2	NA	Unknown Function
Glyma03g28460		NA	Unknown Function
Glyma03g28470		PF05834	Lycopene cyclase protein
	Leaf 1 & 2		NADP oxidoreductase coenzyme
Glyma03g28480		PF03807	F420-dependent
Glyma03g28490	Leaf 1 & 2	PF08241	Methyltransferase domain
Glyma03g28500		PF07719	Tetratricopeptide repeat
	Leaf 1		RNA polymerase Rpb5, N-terminal
Glyma03g28510		PF03871	domain
Glyma03g28520		PF00462	Glutaredoxin
			Cys/Met metabolism PLP-
Glyma03g28530		PF01053	dependent enzyme
			Plant protein 1589 of unknown
Glyma03g28550		PF09713	function (A_thal_3526)
	Leaf 1 & 2		Response regulator receiver
Glyma03g28570		PF00072	domain
Glyma03g28580		PF03088	Strictosidine synthase
Glyma03g28590	Leaf 1	PF03088	Strictosidine synthase
			haloacid dehalogenase-like
Glyma03g28600		PF00702	hydrolase
	Root 1 & 2		<b>Helix-loop-helix DNA-binding</b>
<b>Glyma03g28610</b>		<b>PF00010</b>	<b>domain</b>
	Root 1 & 2		<b>Helix-loop-helix DNA-binding</b>
<b>Glyma03g28630</b>		<b>PF00010</b>	<b>domain</b>
Glyma03g28640		NA	Unknown Function



**Table 3.** RNA-seq and qRT-PCR Expression data

Gene expression analysis of the known strategy I response genes using RNA-seq and qRT-PCR. Expression is reported by fold change between sufficient and insufficient iron conditions.

Gene Name	Annotation	Comparison	DE in RNA-seq Set 1 (Clark)	DE in RNA- seq Set 2 (Clark)	DE in qRT (Avg)
Glyma03g28610	BHLH038	CRSvCRD	3.71*	4.47*	2.99*
Glyma03g28630	BHLH038	CRSvCRD	6.24*	4.11*	2.26*
Glyma12g30240	FIT	CRSvCRD	1.38	3.61*	5.93*
Glyma07g07380	FRO2	CRSvCRD	1.61	3.54*	3.97*
Glyma07g34930	IRT	CRSvCRD	2.22*	3.21*	3.26*
Glyma03g28610	BHLH038	ICRSvICRD	-	-	3.28*
Glyma03g28630	BHLH038	ICRSvICRD	-	-	3.09*
Glyma12g30240	FIT	ICRSvICRD	-	-	0.77
Glyma07g07380	FRO2	ICRSvICRD	-	-	0.70
Glyma07g34930	IRT	ICRSvICRD	-	-	2.11*

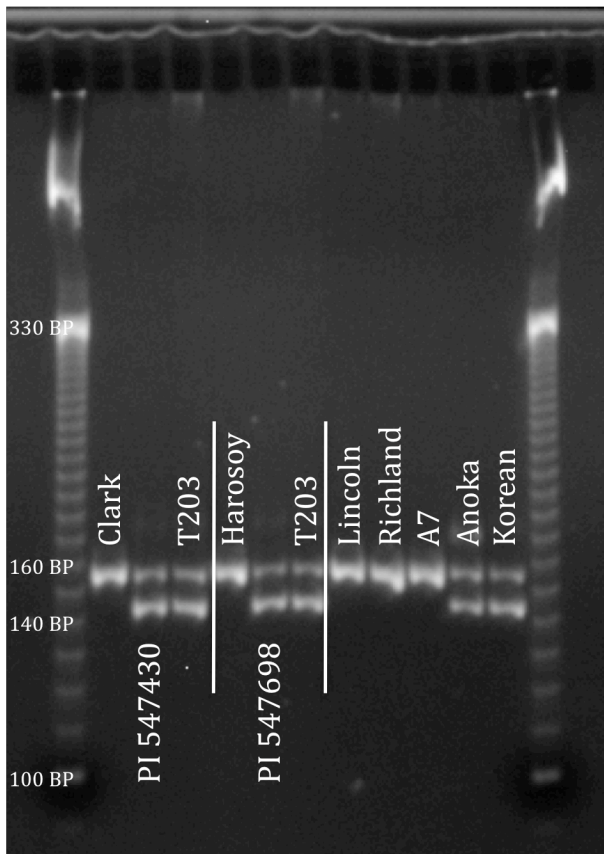
\*Greater than 2-fold expression change

CRS = Clark Root Sufficient

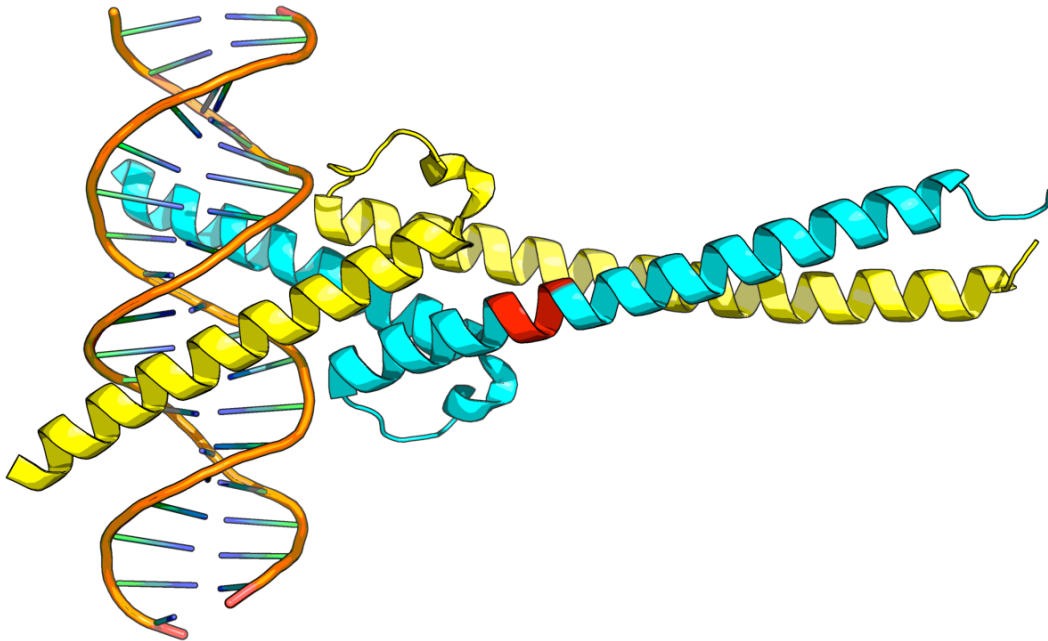
CRD = Clark Root Deficient

ICRS = PI547430 Root Sufficient

ICRD = PI547430 Root Deficient



**Figure 5.** PCR amplification of Glyma03g28610 with primers flanking the 12bp deletion. Due to the high similarity between Glyma03g28610 and Glyma03g26830, the deletion is indicated by the double banding at 143 and 155bps. The deletion is only present in the inefficient lines.



**Figure 6.** Cartoon of the FIT / bHLH038 heterodimer bound to DNA. The protein model depicts the interaction of the strategy I response transcription factor FIT (yellow) with it's heterodimer bHLH038, Glyma03g28610 (blue). The 12 bp deletion is indicated in red.

## Chapter 5. General Conclusions

### Conclusions

Iron deficiency chlorosis is a major soybean nutritional concern first recognized in the early 1940's (Weiss, 1943). Breeding for iron-efficiency revitalized in the 1980's -> 1990's (Cianzio, 1979; Diers et al., 1992; Lin et al., 1997). Researchers identified iron-efficient QTL in various populations uncovering multiple mechanisms of IDC resistance; (1) single major gene with modifiers and (2) a quantitative trait controlled by multiple genes (Cianzio, 1979; Lin et al., 1997; Lin et al., 1998). Since the turn of the century, specific iron response genes have been delineated in model organisms such as *A. thaliana* and *L. esculentum* (Vert et al., 2002; Connolly et al., 2003; Yuan et al., 2008). Using knowledge gained from these other organisms we can better identify underlying genes facilitating the multiple response mechanisms. The research presented in this dissertation was designed to delineate the underlying genetic differences between iron-efficient and iron-inefficient soybean lines.

Previous studies have utilized soybean NILs as a tool to identify genetic differences between iron-efficient and iron-inefficient varieties (O'Rourke et al., 2007; O'Rourke et al., 2007; O'Rourke et al., 2009). The best way to utilize NILs is to define the donor parent introgression sites in the recurrent parent. Chapter 3 of this dissertation identifies a novel method for that type of identification. Through the use of next-generation sequencing and SNPs identified between the parents and NIL derived from them, we were able to fine map introgression boundaries. The study

identified donor parent introgressions on six of the twenty soybean chromosomes. Utilizing the previous QTL studies as a scaffold we aligned the newly identified introgressions with the QTL. Of the six chromosomes with donor parent introgressions, only the introgression on the long arm of chromosome 3 co-localized with any QTL. The QTL on chromosome 3 was identified in the Anoka x A7 population thought to contain a single major gene controlling IDC response (Lin et al., 1997).

In chapter 4 we took the NIL mapping a step further with a second NIL created with the same donor parent used by Bernard (1975). The donor parent introgressions in this new NIL were coincident with iron QTL only on chromosome 3. The two chromosome 3 introgressions overlapped on the proximal side of the QTL. Through Sub-NIL mapping and phenotyping we identified a narrow 250 kb region on chromosome 3 containing recurrent parent alleles required for iron-efficiency. Two homologs of *A. thaliana* bHLH038 were present within this region and were the only genes in the region differentially expressed in the roots 24 hours after being exposed to iron stress. Genomic sequence analysis uncovered a 12 bp deletion in one of the two candidate genes, Glyma03g28610. The 12 bp removes one predicted heterodimerization site and shifts confirmation of two others in the predicted model with the important strategy I transcription factor, FIT. Through PCR amplification we showed that the deletion was only present in the inefficient lines in both of the NILs and the Anoka x A7 population. We hypothesize the deletion disrupts binding with

FIT resulting in the heterodimer's reduced ability to activate FRO and IRT during iron stress.

The identification of the overlapping NIL introgressions in the Clark and Harosoy backgrounds in addition to the Anoka x A7 major QTL suggests that alleles in this region are required for iron-efficiency (Lin et al., 1997). However, it has been shown that there are multiple mechanisms for iron stress response and the QTL identified on chromosome 3 may not be universally applicable. In the quest for agronomically important traits it is paramount to utilize the appropriate tools and resources. The NILs used in this study were the ideal tool for delineating the Anoka x A7 single gene hypothesis. However, in another population their usability may be diminished.

### Future Research

This research has identified a strong candidate underlying the iron stress response difference between Fe-efficient and Fe-inefficient NILs and the Anoka x A7 population. To further advance the knowledge about this strategy I system, this study could be followed up with a knockout or knockdown of Glyma03g28610 gene in the efficient soybean NIL Clark. The knockout / knockdown could be accomplished using either virus induced gene silencing (VIGS) or agrobacterium transformation. The expected results of this study would be that Clark would no longer be able to mount a sufficient iron stress response due to the inability to activate FRO and IRT. A parallel study could also be performed using

overexpression the wildtype Glyma03g28610 in the inefficient lines to restore the iron-efficiency phenotype.

In this research, the candidate gene and the identified mutation were only tested in three populations. It would be prudent to screen multiple populations for this mutation to determine the agronomic value of using these tools in a breeding program. Rather than utilizing multiple bi-parental populations, association mapping would be a useful alternative in the determination of the potential application of this mutation.

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## Appendix

### Supplemental Table S1. SNPs identified by Method 1 on the single-library comparison

This table contains 172 SNPs identified using Method 1 on the single-library RNA-Seq comparison between the Clark/IsoClark 10-day root experimental data. The file contains a unique SNP identifier that includes the genomic location. The sequence surrounding each SNP based on the soybean Glyma1.01 genome assembly is also provided.

SNP	[Clark/IsoClark]
>Gm01_5 3617124	TATTGATACAAAGTTATATTTTTTCATTTCAATTCAAACCTAGCGAACTATGCATCGGTTCCAA AGTTCACATTGGCCCCGACGCAGGCACGCAGCTCAGTTAGCAACAGCGTTTTTTTTTTTTTA TCACTTCATTCCCAACCCTAGCGAACTATCAATCGCTCCCAAAGTTCACATTGGC[G/C]CCC GACGCAGTCTCAGTCCGTACCAAGCTTAGTCCAGCTCCACTGTCCGTAGGGCTCGGAA AGCTCCGCATTTGGCCATTTCGCGCTCTGTAAGTTTCCCTTCTCTAATTCTCTCTCTCTTT CATCAAGCTTTGTTTTGCTTTGGACGTTTTCTCTTTGACCTAAGATCCAAAGT
>Gm02_5 687687	TGACCTACTTAGATTCTGGCCTGATCTGAGAACTGATAATTTTATAGAGAGCAAGTCATTAT GGAGGGACCAATGGAAAAAATTTGGATCCTGCTGTTTTATGATGCCAGATGACTATTTTGTT TATGCACTCAACAACAGAAAGAGATACGACCTAAAGAGAATCTTGACAAGTGCAGG[C/T]AA ATGTACGTATATAAAATGAAATTCAAATTCATGTTTCACACACGAAAAGTTAAGGAAAAAAGA ATAATACTTGTCATGTTTTGAGCTTGAGTTGTTCTTGTTACAATTGGTATCCACTTGATATA TAGTGAGGAATGAGTTTCATACTAAGTTTATATAAAATTTATTCATGAAAG
>Gm02_4 2350182	CGTGGGTGAAAGCGACCACATCGAAGAGCTTTGCCAAGAGCTTGGCGTCGTTGAGGTGCG CATCGACGATGAAGACGTCATGCTTGGCGAGGAGGGACTTGCGGGCTTTTTTCAGAGACG GATCGTAGTAGTCGTTGAAGTTGTCGAGTCCGACGACGCCGTCTCCGCGTCGTTTTAACG[ C/A]GAGGGAGACGTGGGAGCCGACGAAGCCGGCAGCGCCGGTGACGAGGACGGACATG CCGCTTGACGGTGGATCTGGGCGGAGGTGCGGACCTGCTTCTCCCACTGGATCCCGCCC CAGGAGGCGGAGAGGTAGCGGCTGCCGGAATCAACGAAGCCTTGAAGCTGAGATATGA AGCGG
>Gm03_3 6460374	GACATTGCAAATTAATAATTTTCCATACAAGTTTCTCAATCCTAATATAGTGATAGTGATAAA TAGAACTAAGACAACAAATTAGTACTAGGCTCTGTCAATCTTCATAGGCCAATACTTTTACC ACTCTTTAATAACTTATGACCCGATATTTGAAGGTGAGGATAACAATTTGTAACAA[G/C]AAAA TTACAGGAAATACGGATTTTATTACATGCACGTAAGAAACACATCAAATTTCCACTCTCAAG AATCCAAAGAAAAGATAACAATGCATGCTCTCAATCCTCAACATGCTAATAACATCTTGC CAATGCAACTTCCTAAGGATGAATATTTAGACAAAAAAAACATTAAAGAG
>Gm03_3 6554101	ACCCATGTCTAGATGCACTATGGGCATCAGTGACATCAATCGCTCTTTGATGTGAACTCTC GTCCACTCTACAACCAACCAATTATCATAGGTGAAGTCATTTTGTACATGAAAGTTAAAAAATA AGATGTTAGGCCGATTTTATTTAATGTAATTGTGTGTAATAAAATACATATAAATG[C/T]AAAT ACATTTAATATACTTTTTGTTATTCCTTAAAAATTTATTGAAAGATCTAATATTAGACCTAACA TATTCATTTAAGATGTCATCAACAATGATTTAATTAATTTAAATATAGATAAAATTGAGAATT TTTTAAATAGAAGGATCAAATATATTTATTTTAAATAGAGAGACC
>Gm03_3 6559857	TGTTTTCTGCAGGTGAAAAAACTCATAGATTAGAGTCCGAAATTCTAACTGAGAAGCTTTTG TCAATATATGAGAAGCAAAGGATTTTCTAATCAAACGATCTTAAGTGCTCTGATGTTAAATTT GACACCTAAAAGAGTCTTCACATATTTGTCATGCATGAATGGCAAGTTCGGAAAT[T/C]AGAA TGGAGTAATCACATCACATGCCGCTTCTATATACATATGATTTGGTAAAAAAGCAAGTGGCC GGNAGGAAGGTGCATGGATTTTAGGAGGAGTTAAAGAAAAGAATATTGTATAAACATAAAT GTTACAATCAGCCTCACNTAGCAAAGTGTGACACCCTCTTTTCTCACAAATA
>Gm03_3 6559926	ATGAGAAGCAAAGGATTTTCTAATCAAACGATCTTAAGTGCTCTGATGTTAAATTTGACACC TAAAGAGTCTTCACATATTTGTCATGCATGAATGGCAAGTTCGGAAATNAGAATGGAGTAA TCACATCACATGCCGCTTCTATATACATATGATTTGGTAAAAAAGCAAGTGGCCGG[C/G]AG GAAGGTGCATGGATTTTAGGAGGAGTTAAAGAAAAGAATATTGTATAAACATAAATGTTACA ATCAGCCTCACNTAGCAAAGTGTGACACCCTCTTTTCTCACAAATATATGTACTAATAATGA AAGGAATAGAAAGTTAGAATTAATGAAAAGTTTTTAAACACATTTAAATAAAA

>Gm03_3 6560002	CATATTTGTCATGCATGAATGGCAAGTTCGGAATNAGAATGGAGTAATCACATCACATGCC GCTTCTATATACATATGATTTGGTAAAAAAGCAAGTGGCCGGNAGGAAGGTGCATGGATTT TAGGAGGAGTTAAAGAAAAAGAATATTGTATAAACATAAATGTTACAATCAGCCTCAC[G/C]TA GCAAAGTGTGACACCCTCTTTTCTCACAAATATATGTACTAATAATGAAAGGAATAGAAAAGT TAGAATTAATGAAAAGTTTTTAAACACATTTAAATAAAAGCATTTCAAAAAGGATAAAAAGGTT TACATTCGTTTTTCTAGCATTCTAATAAAACTTATTTAAATAAATAATAAAA
>Gm03_3 6952394	AAAAGAAATAAAAAAATAAGATCTGTTTTTTTTTTTTTTTGAGAAAGATCATTTTTTATTATT ATTGTTTTGAATTTTGAGAATTATGTAATGATGAGAGGTTGATGTTGTGAAAAAAAAAAATGG AAATCCAGGGAGGAGGCGGTGGTGGAGGTGGCAAGAGTGGAAAGACAATGTC[C/A]ACCA ACAGCAAGGAGAATTTGTCATAGTTTTTCTTGTTACACTTTGGTACTCTTCAAACATTGGTG TGATCCTTCTGAACAAGTATCTGCTCTCAAACATATGGATTCAAGTTCCTCAATCTTCTCACA ATGTGCCACATGTCGGCCTGTGCTGTCTCAGCTATGTCTCCATTGTGTT
>Gm03_3 6959955	AACGAATTGCAAAAGTTTTGGAAGTGTCTCAAATAAAAATGCTCCTCAAACAGCTGGTTCT GACATAAACAGATATAAACTAGTTATGTTCAAGTGCAGCAGAGAGGACTGTTGTTGGCATATTT GCCTTCAATCATTTCTTTACTCAACACCCATATATGTTTTGTCATGCCTGGAAATT[T/C]AGTA TGGTATCAAGTCTTCTCAAACACTACATAGAATTAGAATGCGGTAACGAATGGCAGCATTAAA AGCACACACTACACTAGCAGAAGAAATCAGCAAAGCAAGCCGGGTAATGCCCAAATGTAAA AAAAGTAGCTTGTTAGGCAATTTCATGTCAATGTTTCATCTACTTGTTTTTCG
>Gm03_3 6996777	CATATATTTGCATTCACTTAACGTAGTCTTGATACCCCAATCCACTAACAATATTTTTTTATA TTATTATTATTATCATCATCATCAAAATGCCTGGCTTGTAATGAAATGGATTGGACCAAAC TATCTAGTGACGTGCGCAATGAAATCATCCAGTTCGCGCGAGAAACACCCCT[C/C]TCATC CTTGGACCTGCGAATAGCTTTTTCAGTTTCATTGCCCTTGTCTCATCTCATCACCTTCTTT TGTTGCCATCAATCTCCTCACGGCATTTCTCAACATCTGATGATGTCAACCACTCGTCCCTGT GGTCCCAGTCCTTCACAACCACCCCAACTCTGAGCACTTCTGTTACCAA
>Gm03_3 6997184	GCACCCCCATTGTCATGCTCTCCATGCAAGAGTTCATCCACAGTGACTCATAAACCCACC AGTTGAACTGTGACTTAGAATCTCCAATTGGGGTGCCAGTCTCTCACAACCAACCCCGTG CCTTTCACCTCTCTCTTCAAACCCCTTTGGAAGCTCAGCAGTTCTCACGCCATCCTCAT[G/C]N AACACATCTCCCTTGTCAGCATCCCTCACCACCCATATGAACTTTTGCTTGCTTTTTTCCAA CCCGTTTGCAACCTCCTTGATTTGCTCCTCAGAGAAGCATGTTGTTGTCCCAAACGACACAT ACAAAACCTGACCCTGCTTCTTGCTTGCAAGCCACTCAACACTGAAGTGCCTTG
>Gm03_3 6997185	CACCCCCATTGTCATGCTCTCCATGCAAGAGTTCATCCACAGTGACTCATAAACCCACCA GTTGAACTGTGACTTAGAATCTCCAATTGGGGTGCCAGTCTCTCACAACCAACCCCGTGC CTTTCACCTCTCTCTTCAAACCCCTTTGGAAGCTCAGCAGTTCTCACGCCATCCTCATN[G/A]A ACACATCTCCCTTGTCAGCATCCCTCACCACCCATATGAACTTTTGCTTGCTTTTTTCCAAC CCGTTTGCAACCTCCTTGATTTGCTCCTCAGAGAAGCATGTTGTTGTCCCAAACGACACATA CAAAACTGACCCTGCTTCTTGCTTGCAAGCCACTCAACACTGAAGTGCCTTGT
>Gm03_3 7024958	GAAATGAAGAATGGGGTTTAGAGTTAGTCAAGACAAGTGGTGTGGTGTCTATGTTATGAC TGCACACATGTGAAGTGAAGTAGAGACTATTCAGTCCACAGCAGCTGTTTCTAGTGTGTGT GTCATTGCATTCTCATCCTTTTCTCTTTTTTTCACGCCTTAATTTCTTTCTCTCTTT[C/G]TC CCTCTTCTCTCTGGAATTTGGAGCATCAGCCAGCACTCTATGGATTCTCTGATTGGTAATT GGCCATCCTACGATCCTCACAACCTTCAGTCAGCTTCGACCTCCGATCCTTCTAGTTCCTCT GTAAGTTGCTGTTGTTGTTGTCTATGAAATTGATAATCCTGGTAATAATTACT
>Gm03_3 7027198	GTTAAGCAACCTTTTTGTAGAAGGGTCTTCTATAATAACTCTAGTTCATTTTCTCCAATGTT ATTTTCTCAAGTTGCTTAACTTTTGTTTTTAAATTCGGTAGACTCAATGGTGTCTTCTAATTC AAAGTTTGCTAACCTGTTGAAGTTCCAACAGTGATAAGTACTGAAGCCAAAAA[C/T]ATCCTC ATGAGACATATTTATCAGCATTCTGAGCAGAAGGTTAGTGACTTGATGGTTAAAGAGCATGT GTTTTGGTGCAGTTAGGTATGCATATGCTTGATGCTCATAACCTTTTGTGTTTTTCAAGTTGAA TCCAAAAAGAGCTGCATCTGATAACCTTCTTTACCAGAGCATGGATG
>Gm03_3 7144398	TCCTTAAAAAGTTGACACCAAACTGAAAAAGCATTATCTATGTTGCAAATGTTTAGGGCTT TCGTAATAATTGCAACAAGCAGATGGGAGTTTTTCAATGACAGGTTCAAGAAAGGTGAAAGA GAACTTCTACATGAGATTCGGCGAAGAAAAGCTTGACCAGCAAGCAACAACCAA[G/T]G CACCAAACCAAGCAACACTCCAAGACTCTGATGAAGATCAAAGGTCTCATCAATCTCATCA TCTTCTGGTTACACTACCTTGTGGATGAAAAACAAGCGACTCAAGAAGGAGAATGGGGTGT TGAACCTCTGAGCTCACAAGTATGAAAAGGAAGTGCAAGGAACCTTCTTGATTGGT
>Gm03_3 7165409	AAGATTCACACGTACCATCTCGAAACAGTTGTTCTGATAACTGTTCTCTGAGAAGCTCCATCA CAGACAGGTGTACCAAGAGATTAGTGTTTTTACTCTTTCTTATTGTCAAGCGACGCCCCAAT TAAGCCTATCATTTCTCGTGTCAAGCTAATCTCCGCAATGGATGTATTTCTGATGTT/GIGAC

	GGAGTCAGTGCCCCCAATGCAAATTTCAAGGTCATTAAGTAAGCTTCTAGGATGAGAAGGG ACATCAAGAAGTCTTTCTTGAATAGCTTGGCAGAGGTTTTGCAAGTCTGCAGAACTCACTTC AGTAGGTTTTGACCTGGAAACAGGGAAACAGAGAATTACATATTTGAAGATGCT
>Gm03_3 7827399	GTCGAGTGATCCATACCCGCTAGGTTTTACACATTCTTTCCCTGAAACAAACACCACCCCT TTTAGGTCTTCTTCTAATCCTCCCTCTCTCTAACTTTCACACGCAGCGGCTTCTTCTCCGTT TTCCTGAATCGGTGTGAACAATGGCGACGTACGCTGCGATGAAACCCACCAAGCC[A/C]GG GTTGGAGGAGTCCCAGGAACAGATCCATAAGATAAGGATCACCTTTCTTCCAAGCACGTC AAAAACCTCGAGAAGGGTATGCACCTATTTATTCTACTTTTTGTTTTAAGGTTGTTGCTATC CTTAGCCTTATGTTTCTAGTTTTGTGTTACTGTTAGTTTTGTGCGGACTTGGTT
>Gm03_3 7828684	CAACACATGGGATAGATTTGAACTTCGTGTCCACAAGAGAGTGATTGACCTCTACAGTTCC CCAGATGTGGTTAAGCAGATTACCTCTATCACAATCGAACCTGGTGTGGAAGTTGAGGTGA CCATTGCAGATGCTTGATCTATATCCAAATTTTAACTGTTATGGTTTCCTAGTTTTTC[G/T]TC TCCTCTTATTCTGTATTTTATAAGTTTTGATTGCGGGACCGAGTTGAATTGTTTTCTTTTGT TACGGTTGTAGACGGTTGATGAATACCTGAGAAATGTCATNTTTATATATTCTAGAATCCTA CATATATTTGTTTTGAAATTATTACGAATCATTTCTAATGGACTCAGCTAA
>Gm03_3 7828791	TGGAAGTTGAGGTGACCATTGCAGATGCTTGATCTATATCCAAATTTTAACTGTTATGGTTT CCTAGTTTTTCNTCTCCTCTTATTCTGTATTTTATAAGTTTTGATTGCGGGACCGAGTTGAAT TGTTTTCTTTTGTGTTTACGGTTGTAGACGGTTGATGAATACCTGAGAAATGTCAT[T/C]TTTAT ATATTCTAGAATCCTACATATATTTGTTTTGAAATTATTACGAATCATTTCTAATGGACTCAGC TAAGATTATATCTTTTCTTTTCTTTTATCATGTTCTGCAAATTTTTAAACCTGAAACTGATTG TGAAGTAGATAAGTTGGAAAGGTGTTTTTGCAAATTTGTACAATTT
>Gm03_3 7832228	CCTTGTGTTTCATGAAACTGCCTTTTGTGTTTTAGGTACCAACACATGGGACAGATTTGAACTT CGTGTGCACAAGAGGGTGATTGACCTCTACAGTTCCCAGATGTGGTTAAGCAGATTACCT CTATCACGATTGAACCTGGTGTGGAGGTTGAGGTGACCATTGCAGATGCTTGATCTG[G/T]A TCCAAATTTTAACTGTTATGCTTCTAGTTTTTTTTCCCTCTCTTTTTGTATTTTGAAGTTTT GATTGTGGGACTGAGTTGGATTGTTTTCTCTAGTTTTACGGTTGTAGACATTGATGAATACC CGAGAAATGTCATCTTAATATATTCTAGAGTGTTACATATTTTGTGTTGAAAT
>Gm03_3 7863675	CTTTGATCTAATGTTTGCAAGAGCATCAGTCCCGCTAACGGTTGCAGCATCCGAGTGCCAC GCTTCGGGAAACGCCTCATCAGCTTGTGTACCCATCCAAACAGCCAGCATCAGCTCCCCCT TCGCCTTGTACCCCTTCTGTCTCCAGCCTATACCACTGCGGTGCCAAGGGACTGTC[T/C] GGTGGAACACGCTTTGGGATCTCATTGAGGTCAAATAAGACACGACCAATGAAGTCATCTT TCACGACATCCTTGTCTTTCACAGTGACCTCCAGTATGGAAGCCTGAATGCGGTCTTTGGA GAAAGCAAAAACCTGGTTCCATTCCGGATTAGACTTCTTGTCAAGTGCCGAGTGGT
>Gm03_3 8065066	TGTGGAACCACCACTCAAGACTATGTTACCATAGAGATCCTTCCTGATGTGACATCACACT TCATGATAGAGTTATATGTTGTCTCATGGATACCAGGAGATTCCATCCCAATCATGGATGGC TGGAAGAGAACTTCAGGGCACCGGAATCGTTCAGCACCGATCGTAATCACCTGCC[C/A/G]T CAGGTAGCTCATAGCTCTTCTCAACTGCTGAGCTGGTCTTGGCAGTTTCCAACCTCCTGCTC ATAATCAAGAGCAATGTAGGCCAGTTTCTCCTTCATGTCCCTCACAATTTCCCGCTCCGCAG ATGTGGTGAAAGAGTAACCACGCTCAGTCAAGATTTTCATCAGGGCATCAGTGAG
>Gm03_3 8083417	GTTTTCTTTTCTCTTCCATCCATGGATTCTCTGAGTTCCATCAACGGCGATTTCGGCTTCGA CGGCGGCACGGATCGCAGATTGCGGTTTTGCGGCCAAGCCTCCTTCCAACAACCTCACAC GCCGATAGACATTCCGGCGCACGGCCACTACGACCACCACCACTACTGGTCCCCGCGT[G/ A]ACGATAAACCTCCGCCGCTCTCTCCAAATATCCTCTCTCTCCTCCTTCGTTTTCTCCGT TTCCGAAGCGTTAGGTCCGGCCACAGGTACATGAAGAGGCTCTTCTCATGATCTCGCTC AATGTCGCGTACTCCACCGCCGAATTGCTCACTGGACTCTTACCAGGTCGCGTAGGT
>Gm03_3 8117453	GTACATGAAATTATAATAACCGTTTTCTGCACTTGTAATTATTGGCCTTACTTGCTATT CTCATGAGCTACAGCAGCAATGTGGTTGCTTGTGTTTTGCCAGCAAACCTGCTTATGGAACGC GCTTTACCATGCCAAGCTTCATCCTCTTCTTCGTCGACGTCATTATTATCCCTGG[C/T]TTC TTCTCCTCCTTCTCCCCAAAGGGCTCAGNATCATCAACACACTCAATTTGCTTCTCGAGAGA ACAAAAAGAAGGGGAAGAAGAAGAAGGGTGAGGCACGTGGTGCACGTGCACCTCCTTTGG TGCAGTTGATGCAATAATGGAACGGCTTCTAACACTTGTGTCATGCTTCTGAGTCT
>Gm03_3 8117485	TTGTACTTAATTATTGGCCTTACTTGCTATTCTCATGAGCTACAGCAGCAATGTGGTTGCTT GTTTGCCCCAGCAAACCTGCTTATGGAACGCGCTTTCACCATGCCAAGCTTCATCCTCTTCTT CGTCGACGTCATTATTATCCCTGGNTTCTTCTCCTCCTTCTCCCCAAAGGGCTCAG[C/T]ATC ATCAACACACTCAATTTGCTTCTCGAGAGAACAAAAAGAAGGGGAAGAAGAAGAAGGGTGA GGCACGTGGTGCACGTGCACCTCCTTTGGTGCAGTTGATGCAATAATGGAACGGCTTCTAA CACTTGTGTCATGCTTCTGAGTCTTCCCTTCCCCAAGTGATGCTCACACAGCGAGT

>Gm03_3 8132996	TGACAATTTACCTATAGAAAAATGGTTGATTTAATTACATAATTACTATGATGCTTATTAAGTA GTGGTTTTGTTTTGAGTGGAAAAAGGTGCACATAGTTTTGGTCGTGCGCAGTGATAGATTTTTTC AACCAGAGACTGTTCAACTTCAGCGGCACAGGTAGCCCTGATCCCACGCTGAAT[A/G]CAA CCTATTTAGCGACACTTCAGCAAAACTGTCCACAAAACGGAAGCGGCAACACTCTGAACAA CCTCGACCCCTTCGTCTCCGGACACCTTCGACAACAACACTACTTCCAAAACCTTCTCAGCAATC AGGGCCTTCTCCAAACAGACCAAGAGCTCTTTTCCACCAACGGCGCGCCGACACA
>Gm03_3 8136913	TGTATTGGATCTGGGGTCCCTCATGTTTGCGACGAAATTAGGGTTTACTCCCAATTTGCCCC CAAATTTTGGGGGTTTTGTTTTCTGATTGTTCAATTTTCATCACCCGACCCCCCGTAGAACCT GGGTTTCTTCGATTTGTGTTAAACTCGAGCTTTTTGGTGAAAAAATACTCGCTTTC[C/A]TTT TCCCTTTGTTGTTCCCTTTCCCTGCAATTGAAATTTGTAATAAAGGAAGCTTCTTTCTAGTTT TGTGATTGAAGCTGCGATGGAGGAAGAGACCTCCGATGCAATGAACCTTGATTGAATCTG GGCCCAGGCCCAGAACCCGAAACTGGGCCCATAGCAATGAAGCTGTGAACT
>Gm03_3 8170431	TTAGATATCTTAGTATGGGAAAAATAAACGAAAAATGTGAAATCTTTCAGCTGTGGAAAGAT GGAGAGATGAAAGAAGAAGTCATTGGAGGCCATAAGGCCTGGCTAGTTATTGAGGAAGTC AAAGAAATGATCCAAAAGTATTTATGATTCTTCTATTTTATTTTTCTTGCTCCCTCAA[C/T]CC CCCACCCACCTCACAAAAAAGAAAGAAAGAAAGAAAGTTACAATTGAAGAAACTAA TCTTCCAATGTTTTACTTTTTTTCTTTAACATTTTTCATGTTGGCAACCGCATGTAGCTGTG TATCATATATATATTAATGGAACGTGTACCATAGCTGAAAGAAATTTA
>Gm03_3 8173719	CGTCGTTGTCGTTGTCGCCGCCGGAGACGGCGACAACGACGACGGCGCAACCTTCTTGG GTTCCAATTCGTCCTTAACAGGACCTCCTTTGGCCATGCTCTGCACCCAAACCTAACCTTC TCCAAGATCAAGGGCTTCCCCTTGAAACCATGGCATCCTTAATCCTCTTTCCAACCGG[T/C ]CCCTCGTCGTCGACAGTAACCTAACCTCGGGATCCTCGTCGGCGTTCTCGTCGGAGATG TAGGGGATCTCGACGGTGCCGTCGACCTTGAGAAGNGAGGTTCCCAGGGTGTCTTTGGCC TCTCCTGCCAATTGAGAGTGAGGCTGATCTGCTAGCCGGGGATGATCTTCCCCTTTTCG
>Gm03_3 8173815	GCTCTGCACCCAAACCTAACCTTCTCCAAGATCAAGGGCTTCCCCTTGAAACCATGGCA TCCTTAATCCTCTTTCCAACCGGNCCCTCGTCGTCGACAGTAACCTAACCTCGGGATCCT CGTCGGCGTTCTCGTCGGAGATGTAGGGGATCTCGACGGTGCCGTCGACCTTGAGAAG[A/ C]GAGGTTCCCAGGGTGTCTTTGGCCTCTCCCTGCCAATTGAGAGTGAGGCTGATCTCGTA GCCGGGGATGATCTTCCCCTTTTCGGACATTGATGTAGGCCTCGCCGTCGAGGGAGCGAAG CGACGTCGTTTTGAGGAAGAGGTTACCCTCGCCGTCGAGGATAGGGAGGTTAGTGAGGAG
>Gm03_3 8174150	CGAGGATAGGGAGGTTAGTGAGGAGGCTGGTGAAGAAGGTCTTGACCAGTCGAGGCAGTT GGTCTCGGACCAATGCCAGTTGTGGACATTGGTGCCGTCGGGGCGGTCTCCACGATCCA CCGCTTGTCGCCCTCACCGTAACGAGCCATTGCAACTGATGCCTCTTTTCTGGGAGAGTT[C /T]CTTAGTNATTAATAATAATTGGATTGGGGTGAGGGTTATGAACTATGATGATTATTATTAT ATGTGTTGAGTGTGTGTGGAATGGTCTAGAAATGAAAAGGAAGGGTCTAGAATGTTGATGTT GCTTTGGGTACACTTTCTTCCACCAACCAACCAAGCCGAGAAGGTTCCCTAGGGTT
>Gm03_3 8174157	AGGGAGGTTAGTGAGGAGGCTGGTGAAGAAGGTCTTGACCAGTCGAGGCAGTTGGTCTCG GACCAATGCCAGTTGTGGACATTGGTGCCGTCGGGGCGGTCTCCACGATCCACCGCTTG TCGCCCTCACCGTAACGAGCCATTGCAACTGATGCCTCTTTTCTGGGAGAGTTNCTTAGT[G /A]ATTAATAATAATTGGATTGGGGTGAGGGTTATGAACTATGATGATTATTATTATATGTGT GAGTGTGTGTGGAATGGTCTAGAATGAAAAGGAAGGGTCTAGAATGTTGATGTTGCTTTGG GTACACTTTCTTACCAACCAACCAAGCCGAGAAGGTTCCCTAGGGTTTTCTTTT
>Gm03_3 8718231	GATATAAAGAGGCTTACCGTTGATTTTGATTGAGGTTAAGTTTATGAAATACTCATCAGACG GTTGTGGATTGTCCGTGACGACGGTGTCCGCAACGGGGTTTACGATGAGTTGTGTGTAAG TAAGGTCAATTTTGAAGAGAAAAAGAAATGAGGAGGCAGTGAGGCGAATATAGCTGC[G/C ]CCTGTGTTGGCTGATGAAGCTGGCAGGCAGAGAGTGAAAGAACGAGGTGAGGTTAATGAA GTGCTGATTTGGGCCGGGAGGGAGTAGTTGGACCGGCCCAATGAGGCGAGGCCCAATGC ATTGGCGGCGAGGGCCCTGGAGAAGATGGGCCGTGCGCAGGAGAAGATGAAGTCGGAGA T
>Gm03_3 8718672	GACGGGGTTTTCGGGGAAGAGGGCGCAGAGGGAGGAGTTGTTGGAGCATGCGTTGGAGG GGAAGGAGTTGCAGAGAGGGGTGTTGCAGGGGATGTGGTGGGAGGAGGAGGAGGTGTAG GTGGAGTCGCAGAGGACCCAGCTGAGGGAGGAGCCAAGGTGGAGGTGGAGTTTTGTGGG TTG[A/G]AGTGGGGTTTTCAGGAACACGGAGAGAGTGTAGAGTTGGGTGGTGTCTGCTTTT GAGATTGGGGCTATGAGGGGGATTTGAAAAGTGGGTGTGAGGAAGAGGAAGAGTATCGCC AAGTTGAAGTTGCAGAGAGATGGAAGAGGAGGAGGCATTTTGGTTTGCTTGCTGGAGCAA GGAGA
>Gm03_3	TAATTTCTTTTTTTTTTCGCATTTCTATTTTTTCTCCTCATTTCTTGGAACCAACAGTTAG

8942920	ATTAATGACAGTTAGATTGATGAGGTACCTGAATCCTGACATAGTCGGTACGATTCCAACAG AAGGCGTTCCCTAGCATCTGGTCAACGGTCTCTGAAACACCGTCGAGCGCAATG[C/T]CAAT CANCATTGACGTAGAGCACTCGCCGGCATTGTTTCATTCTCTTCGCCGGAGCTCCGTTTCCG ATGGAGAGAAACAAGGAGATCCTCCACGCCGTTACCGACGGGAAGTCGCGCTTGTGTGG AGGACGTGCGTGACCGCCGCCGCCGGCGGATTGTTTCATCACTAAGCCGCCGTCG
>Gm03_3 8942927	TTTTTTTTTTCGCATTTCTATTTTTTCTCCTCATTTCTTGGGAACCAACAGTTAGATTAATG ACAGTTAGATTGATGAGGTACCTGAATCCTGACATAGTCGGTACGATTCCAACAGAAGGCG TTCCCTAGCATCTGGTCAACGGTCTCTGAAACACCGTCGAGCGCAATGNCAATCA[C/G]CAT TGACGTAGAGCACTCGCCGGCATTGTTTCATTCTCTTCGCCGGAGCTCCGTTTCCGATGGAG AGAACAAGGAGATCCTCCACGCCGTTACCGACGGGAAGTCGCGCTTGTGTGGAGGACG TGCGTGACCGCCGCCGCCGGCGGATTGTTTCATCACTAAGCCGCCGTCGACGGCGG
>Gm03_3 9788794	ACTGGGTGAAACGAGCCCGCAAAAAGAGGGGTATATGCGAGTGGAATATTTAGTTTCGAAA CTGTATTATGTTTCACTTATGCGCTGTTTTTTCTTTCTGTTCCGTTTCCTATAAATCCATGCC ACACCCTTCACCTTATTTTCCGCACATCATATTACTCTTTCACATCATTTCTCAA[T/C]CTTCN TTGCTTGCACTCTCATCAGGTCAGTTTCACAACCCCTTCTTCTCCCTCTCTCAATATTGTTTT GTTTTTTCTCTTTGGCATCATTATCTTTCTCATGTGTCACCCCATGCACATGCATCTCGCGA CTCCATCACACCGATTCTCTATTTTCTTTTCATAACAAAATACTGAGA
>Gm03_3 9788799	GTGAAACGAGCCCGCAAAAAGAGGGGTATATGCGAGTGGAATATTTAGTTTCGAAACTGTA TTATGTTTCACTTATGCGCTGTTTTTTCTTTCTGTTCCGTTTCCTATAAATCCATGCCACAC CCTTCACCTTATTTTCCGCACATCATATTACTCTTTCACATCATTTCTCAANCTTC[C/T]TTGC TTGCACTCTCATCAGGTCAGTTTCACAACCCCTTCTTCTCCCTCTCTCAATATTGTTTTGTTT TTTCTCTTTGGCATCATTATCTTTCTCATGTGTCACCCCATGCACATGCATCTCGCGACTCC ATCACACCGATTCTCTATTTTCTTTTCATAACAAAATACTGAGATATAT
>Gm03_3 9790874	TTCAAGTTGCATTCAAGGAAAGGTTTGATGCTCCACCATATGAGCTTGATCCAGCTGGCTCA ACTGTTGCACAAGCAGACTACATTTGGAGGATAATTGTTATGGTGGGAGCACTGCCAGCTG CGTTAACTTACTACTGGAGGATGAAGATGCCGGAACCTGCCCGTTACACCGCTCTAGT[C/T] GCCAAGAACACGAAGCAGGCTGCAGCAGATATGTCTAAGGTTCTGCAGGTTGAGATTCAAG CTGAACCGCAGAAAGAGGAGCAGAAGGCTAACTCATATGGCTTATTTTCAAAGGAGTTCTCT CCGTGCGCATGGACTGCATCTACTTGGTACAGCAAGCACATGGTCTTGCTTGATAT
>Gm03_3 9795164	TCTGCAGATGACAAAGCATGAAGAGGTAATTCATTAAGGTAAAACCTAGTAGATTCTGATA TGAAAATGAATTTAAGTGCTTAAACATTTCAATGTGCACGGTCTCTCAGCACTTCAAGATCC ATTTCTGTGGGGTCAGTAGCCTGGATTTTCATAATGGACACCATCTAGCTCACGTCT[A/G]AA CCTGTCTTTAGTAGTGGCATAGAGCATCTTGGCACGNATCCGAGATGTTGAAGGGGACCTA ATCAATATCATGCAATGAATAACTAAGAGTGATATCTCTGGCAAAAGCATAGAATGAGTAAG TTAAGAAGCATGTTGTTGAGGATACATGAGAAGGCTGGGATAAAAGAAGTTCAA
>Gm03_3 9795203	GTAAACCTAGTAGATTCTGATATGAAATGAATTTAAGTGCTTAAACATTTCAATGTGCACG GTCTCTCAGCACTTCAAGATCCATTTCTGTGGGTCAGTAGCCTGGATTTTCATAATGGACAC CATCTAGCTCACGTCTNAACCTGTCTTTAGTAGTGGCATAGAGCATCTTGGCACG[G/A]ATC CGAGATGTTGAAGGGGACCTAATCAATATCATGCAATGAATAACTAAGAGTGATATCTCTGG CAAAAGCATAGAATGAGTAAGTTAAGAAGCATGTTGTTGAGGATACATGAGAAGGCTGGGA TAAAGAAGTTCAATCCAAGTTTGTCTTTCACAAGAACAAGAGAGGAAGACAA
>Gm03_3 9964048	CGGGCCATTCTGCACCGATCAATGGAAGAATTTCAACAACAAAGCGAATCGCATTACAA TTTCGATGCGGATCGGCTTATGTTAGGGTTTGAGATCGAGAAGAAGAGAATTAGATCATAC GCGGAAGCAGCAAGAAGCTGAGAAGAAATGCAATTGGCAAGCGAAGACTTCGACTTCC[A/T] JGTGAGGAGTGGCTTGAGTGCAACTGAAGTAGAAGTCGTCGTCGTTGGTCGTGGCATT ATTTTAGTTTCCAATAATATCCTTGAATCCACGTTTTAACTATGCCTATCTTCGCAAAATCAG AAAATGTTTTTTTTTTTTTACAATAAAGTTTTAGACCAGGGTACTACTTGCATATT
>Gm03_3 9966061	ATTTTGAGAATATTACATTGACTGCAACACAACCTCAAAATCCATTAGTGCAAAATTAACATTC GTCGCTTCATTCTAATAACGTGAGCCAGATTTTGATGTAAGTGGTTCACAACACAAAAGGCA TAAATGGCTCAATCACTTGACACCAGGCACCAGCACTAGTTTCACAAAATACTGT[C/T]ACG CATGGAACAAAAGACACTTAAAAATTTAGAGCTTCTACATCACTCAATACTTTGGAGGTGGCT GCTGGGTTGGTAATAATCCAAAACGCTTCTTTCAGAAGAACTCTCTGTCTTGAATATTTATCA TCGGGTGAAAACCGGGCTGCAGTTCAGGAAAACCGGTAAGTACAGAAAATG
>Gm03_3 9967974	TATCGTTTATGTAAACCTGAAGATACATGGTCACTGCAAAAAAATTGAAATACATGAAATCAT TGCCAGAAACATAGCTCGTCAGATGGAACAATAATGAGTCTAGAGTTTTTCGGAAACATAC CTGATCTTTGATCGAAGGAAAGAACAGCAATAGAGGTAGAAACCTGCAATGGAGGC[G/C]A GAAAAAGGGTTAAGGTTTTCGGATCCTCTGGTCTATACAAAGTGTGTTTCAACGAGAA

	TGGGCCTGTCATTTAGGTATTGGTTATGGGAGATTTTATTCCTACACCGAGCATATTTCTTC CTGCACCCAGCATATTTATGGAAAACCAAAATACCTTACGTATAGTCTATAAG
>Gm03_3 9986148	TTGTGAAGATGCTGCATGCAATCATTTGTTGAAATTTTTATGGCTGGAGTTTTATTTACTT TCTACTCGTGTGTTCTTAACTTTTTTATTTTTATCTTGTCTCTTAAGTATCATGATCAAGAGAA GGGACGCAATTCGTCCCGATATGGCAGCAGTTCAAGAAGACCCATAATCTCAT[C/T]GTCAA CCAGGCCAAGTTCCTCTGGTGATCATACTGACAGTCGTACTGGCCGGCTAACCTCAAGTGG AAGCCGACAATCTGCCACACATAGAAATATTCAACCTATGCACGAGACAAAACAACCAACTT ACACACGCTCTGGATCCACCAGAGGCAACCGTGATGATCCTCTGCGGAGTT
>Gm03_3 9993121	CACTATTTGAAATATTATAATTAGGTGAGAGGGAAAACGATGATTGGGAAGGAGAAGGAGA ACCTTTGGCGGTGGGTTTTCTTGGTGAAGATGTTTCATGGCGATCGGTGCGTGTGTGCACAGT CGAAGGCAATTTACGGAGAGACACGCAAAACACAACCCAACGATGGTGAGAGAGAAGAAA[G/ A]ATCCTCGCAATTTTCATGTTTTATTTTTTTATTTTTTTTTATAATATATGACTTTCTCATT CGACTATATATAATATTTAAGATTTTCAGTTAACAGAAACCGCATAGGTCTTGGTCGGGTAT TTAAGCATATAGTTGACTTGTCTAATTTACATAAGATGCTTATATGTTTTTG
>Gm03_4 0055679	CATTACCTTAAAGACTATTTAGCCGAGAAGGTAATACTTGAAGGAATAAATTGATGGTTTTAG CCTTATTTTTCTTTTTAATGGCTTAAGCTATGTTTATTTTTCTTCCAAATTTTTATTTTTTGGT TTGACATCTAACCTATACTTTCTATTGGTGGCAGATGTAAGGCACATACCTTT[C/T]GTGGAA AACGTGGTGGCAATATATAGAAGCTTGGATGCCACTTTTTGTGCTAGTTTGTTCAGTCGCG TTCTTCTGGTGAAGCCTCTTTGTTTTGGAGAGCTTTCTGATTGATTGAAGAAAAAAAAGCT ATGTATTACTTGATGCTGATTTGATATTGTCTTTTAAAATTCGTTGAT
>Gm03_4 0056070	AAACAATTCCTTAATTCAACTTCTAGTAGTTTTAAAACTGGCAAGCTTCCTCGGAACCTAACTA ATTTGAATGAGGTTGGGGTTGGCCAACTTTCAAATCCAAGAGTGTGGTGCATGATAAAGC TGAGGAGCTTGAGTTGTCTTGTGGGAGAGGGAAGCCCAATCCAATGCAATCTTCCC[A/T]GA TGGGACTTGGGGGCCGAGAACAGTGTGGAGTTATGGAAATGGATGTGAATCAAGTAACAC AATCAGCACCTGATAGCCCCGCCCTTTTGTGATACCAAAGGTTGAGATAATGACTGCAGTGA TCAAGATAGTGAGCATGTAAGGAAACATACTCTCTTATTTATGGTCTTCTGTTTTGT
>Gm03_4 0131229	TTATATGTTAACTTTACCAAGAACAATAAACCACATAAGCCAACCCAGTCAACCCCTGAA CAACATTTTGCAAACAAGGTGCCCTCAACCTCATTCAAATCCATGCACTCACAATCACAA TGAGTAAAGCAGACTATAAAAAAGAAATTAAGATCACAGAAAAACCAAGAAAAACA[C/T]CTC TTATACTATCAATTAATCTCATCAAAACACTTCTCACTGAGAAGTCAAGCAAAAAACAACTA TATGTGCCAAATGCCAATATTGTCTGCAATTCAACCTAAAACAATATCTGAAACNACTTCAA GAAGCAGAAACATTACATAAACTAAAAACCCAGCTTAAACAATAGCATTG
>Gm03_4 0131350	CAATGAGTAAAGCAGACTATAAAAAAGAAATTAAGATCACAGAAAAACCAAGAAAAACANCT CTTATAACTATCAATTAATCTCATCAAAACACTTCTCACTGAGAAGTCAAGCAAAAAACAACT ATATGTGCCAAATGCCAATATTGTCTGCAATTCAACCTAAAACAATATCTGAAAC[A/C]ACTT CAAGAAGCAGAAACATTACATAAACTAAAAACCCAGCTTAAACAATAGCATTTGCAGCACTT GCAATCCTGGGCCACAGATCAGCACACTCCTTCCCAATAGGACCAAGTATGAGCAGAACCTA CAATTAATAATCTCACCAATCAAACACTAAACATTTACAGAGGAAAAAAA
>Gm03_4 0132046	GATTCATATGACCCAGCATTATCTGCCAAAACGCAACAATAACAATTAGAAAACAGAAA AAAAAATTAACAAAGACAGAGATCAATAATAAGGAAACAGAAGAAAAAGTGAATACCTTCG AAGTACATGTAACACCGTCCTTTGCGGCCACGGCTTGCGCTGACGGACGATGAC[G/A]G CGGGGAGCACCTTTTTACGGAGATCAGGCTTTCCCTTCTTGACGGTGGCCATAACCATGTC ACCCACGCAAGCTGACGGCAACCGGTTGAGCCTTCCCTTGATCCCCTTCACTGATATGATG TAGAGATTCTTCGCGCCGGTGTTGTGCGGCGCAGTTCACCGTCGCGGCCACCGGCAG
>Gm03_4 0154304	TTATTCTAATGTTGGTGACGTGAATTCGTATGCTGGGGATCCGAATTCATCCTACAGCAAG CTCAATTCAGTGCAACTGGTGAGTCCAAGCCAGCTGGTGGAGCGGCTGATTCTAACGAGG CCTCTGCTGGGGTTGGAAGCAGCTGCTGACTCTACTATGGTATCTGACTACAATTC[A/G] TCTGTAAATGNTGGTGTGCGCGTGCAAGTACAAATACTTCTGGACTTGAAAACGGGAATG CGTTGGAGAATGCTGATGGGTGCGCTGATGAGAAACAGCAAGCAGATGGTTATGGTGTGC CTTTTTCTTTTACCTTTTTACTTTCAATGGAATTATCATGTTTTAAATTTTATTTTTG
>Gm03_4 0154315	TTGGTGACGTGAATTCGTATGCTGGGGATCCGAATTCATCCTACAGCAAGCTCAATTCAG TGCAACTGGTGAGTCCAAGCCAGCTGGTGGAGCGGCTGATTCTAACGAGGCCTCTGCTGG GGTTGGAAGCACAGCTGCTGACTCTACTATGGTATCTGACTACAATTCNTCTGTAAATG[G/A] JTGTTGTTGCCGGTGCAAGTACAAATACTTCTGGACTTGAAAACGGGAATGCGTTGGAGAA TGCTGATGGGTGCGCTGATGAGAAACAGCAAGCAGATGGTTATGGTGTGCCTTTTTTCTTT ACCTTTTTACTTTCAATGGAATTATCATGTTTTAAATTTTATTTTTGGACAATAAATT
>Gm03_4	ACTACTTGAGGTCTGTCTTCACTATTACTTTCTTGCTGCTAAATTCATCATATTATGTA

0160427	GTTGGTGATTCTTCTCTTTGTATTCTAAGTGTTCTTCATCTCTAATGAGATTCTTTAATTCAA TGTTACATGCAGGCTATATTGCATTTTGAGGCTATTCAACCCTTGCCAAAGCG[C/T]GTTGAT ATTGACTTTCTGGAGTCATGGGTTGTCAAATTCATAATGCCTAACTCAGAGAGTCCTGGTGT AGCAAGTGCAACAGAGCGTGAGGAACCTTTCTAGTATTTTCTGGAGGTAATTCCCTGTGTCAGA ATGGTGTTCTGGCATGTGTTCTAAAGGAGTTTCCTTAAGAAATGTGTTT
>Gm03_4 0172340	CAGACTACGGTTTGGAGGAAGAAGCCAGGGAATATGTCAACTCAGTTCATATAGATGATGA TCCAGTGGACGAGTATAGTCTTCCTGAGCACCAGCAGCAGCTACAAGAAGAAGCTTGAACCC GAAATTGTGGAGGAGGAACTCCCGTACAGGAGGCATCTCCACCAATTCATAGCATTG[C/G] JACACACTGTCCAAGAACCCCTGTTGCTCTTGTGGAAGAGTCCTTTGAGGAGCCTCCTAAG AAAACATATGCATCTATTGTATGTAATACTTTCTATACATGTTCTTTTTTTTTTGGCAGACTAC CTTTTTTGTGTGTAGTTATGTACTATCCATAAAAAAGGTACCACATGACTGGA
>Gm03_4 0179701	CCACCATTGAGCAGCACTGAACCCATCTTCACTCTCACTTTCCATTTCTTCATCTTCTCTTCT TCACTCAAGCCATTGTGCTTGGCAGTAGAGAAAGAAGGAATGGTCTCCAGCACCAGCTCC TGTGTCTAGCTGAAAACCTCTTGGAAAGGCTGAAAACCTCAAACCTTTCTTCATTG[T/C]TGC AAGAAAAGAAGGCCAAGTCTTGCAAGTTCCCATCAGATTGGACTCTTCTGATCAAGCCTTT GTTTTGCCTTTCAAGCTCACCAATTGAGGGAGAGATTGGAGAAGAACCACATGCAGAGGAC CCAGTTTGGTGGAAGTAAGTTTGTGATAATGTTGTGGGACAGTGGTTGGTGGGA
>Gm03_4 0211426	TGGCGTACCAGCCGGTGTGCGGAGAAGACGCCGTTGACGTTTTCCAACGGAGGACCGAGG CCGGCGTTGGTAGTGAACCTGCACATGTTGGTCCAAAGGGATAAGCTTCTGCAATGCTTCA ACATGTCCTCGTTGAAGCGCGAGGGAAGGTCTGGACGAAGATGTAGCGTCTCCGCATG[ G/A]GTCGCTCTTGTGTGCGCGGTTCCGAGGGCACGGTCAAGGGGAAAGTGGGTTCCCTT CGGTTCCGAAACTCGCTCAGGGAAGCCGAGGTTTTTACGAGGTTGTTGGTGTGTTGTTGG GATTGGAAAGTGATGGGGACTGCTGGGTCAGGGGCGGTGGGTTGGTGGCGGAGGACGGC GA
>Gm03_4 0371846	CAGAGCTCTTGGAGCAAGAAGTGGTCTCCCTTACCGCATAGATCCATGCTTGGCATCCTTC ATGCAGTAGCATTCTTTACTACCACTGAAGTGAGCCAAATGGTTTAGAATAAATTTTTTGTT AGAGGAAGTGATATAATATTGGACATTTCTTCTACATCTTGTTCATACTTTTCA[T/C]AAAT GTCAATTTAGAATTCAAAGCCAACCTTAACTGCTGGAAAAGCTACCCCTTTCTGTTGATTT GTCACACACTTGCCTCACTATGGTGGTCTATATATAAAGTCCCAAGATACCATTCCATTTT CACAATATTGATGTCTTCATAAAGCATGTGGTCTCGTGATTTAACGTTGA
>Gm03_4 0459321	ATTTTCAATTTGTTATATGATTTGTTGTACTTGTTTTGTTTATTTACTTATATTCATTTTCAT GGACTTTGCAGATTGTTTCTGACTCAATGAGAGCTACAAATAAGAAAATAAATGGAACACCC ATCAAGATGCTCATAGACCAAGAAATGTCCAAAGAAGTTGTTTCCAAGCACA[C/T]CCACC ACCAAATGTAGTTGCAAAATTAATGGGGCTTGAAGCCCTCCACAGGGTGAGCTATCTGTG GAGAGAAGCCATAGAGGAGATTATTTCAACATATGTGTGGTCATTGAGGACACCATTCA ACCACTGGAACCTGGAAGATAGGTTTATGGACAAGGAAATGCTTCATGAAGT
>Gm03_4 0462434	TGGATCCATCATTGAGGGACAAATACGCCAATCTCAATGACAAGGAGCCTCAGCAGCAACT CCACGAGGCCAAGCGAAGGCAGAGGAGGTCCAATCAGAAGCTTGATTTCGATTGTGTGAA TGTCTCACTAATAGAAATTACTGGTTATGGATCAGAGAAGAATTACTTGATGGGTAGTA[G/T] GTTGTGCAGTGGGAGCCACAGCAGGGTCCAAGTCCAGAAGCTGCATCTCCCCCATTGGT GGACCTCATTGTGGCACAGATGAAGGAGTTAATATCTAGTGCTATGAGTTCTGTTTGGGTG GTGGATTGTGGGGACAGTAACAGCCTGGTGGTAGAGAGTGTTGTCAGAAAAGAGGTTG
>Gm03_4 0462640	GTCCAAGTCCCAGAAGCTGCATCTCCCCATTGGTGGACCTCATTGTGGCACAGATGAAG GAGTTAATATCTAGTGCTATGAGTTCTGTTTGGGTGGTGGATTGTGGGGACAGTAACAGCC TGGTGGTAGAGAGTGTTGTCAGAAAAGAGGTTGTGGGCAAAGGGTGGGTTGAGCTTATG[A /G]GATTGGAGATGGATATTTTGGTGAAGGAAGTAGAGGGGAAGCTGCTAGAAGAAGCTTGTG GAGGATGCGGTGGTTGATTTGACAGGCAGGGCCTGAATGGTTGAAAACCTTGACATGGCT TTCCGCTTTGGCTTTGCTCTTTTTTATAATATATATTGTAACCTACTATTGTTGTATTC
>Gm03_4 0585266	TAAAGTGCATAACAATTGAGTATACCATCTAATAATGAAAATAACGCAAAATAACATAATGCTT AAAGCAACGTTTCATATCAGAAGGGGAAAAATAGTTTGAATCCCATCCAACAAAAATCATGCGA TTAAAATCCAATTAAGGAGCATAAAGTAAAATAAGTTTTTAAAACAGGTATCTG[G/A]AGGA AAGCAACTACATTTAGTCCACTTCTCAATCTTGGGGCCAGCACCACCTCCACCAGAAGGA GGAGCAGCATACTCATCCTCTGCTGCACCAGCACCACCCACGTCTGGACCAGCACCACCT TGGTACATCTTGGCAATGATTGGATTGCAGATGCTTTCCAATTCCTTCATTTTG
>Gm03_4 0585499	TCCACCAGAAGGAGGAGCAGCATACTCATCCTCTGCTGCACCAGCACCACCCACGTCTGG ACCAGCACCACCTTGGTACATCTTGGCAATGATTGGATTGCAGATGCTTTCCAATTCCTTCA TTTTGTCCTCAAACCTCATCTGCTTCTGCAAGCTGGTTGCTGTCTAACCCTGGATTGC[T/C]T



	GCTCAATTGCATCCTCAATCTTCTTCTTGTGTCAGCCGGGTCAAGTTTCTCACCATCTTGTCA TCCTTCACGGTGTTCCTCATGTTGTATGCATAGTTTTCCAAAGCGTTTTTGGCCTCAACCTT CTTCTTGTGCTCTTCATCCTCAGACTTGACTTCTCAGCCTCTTGAACCATCTT
>Gm03_4 0586108	GTCAGAGTATGTTGAGAAAACCTGTTCCCTTCTTGTGGAATTGTAGTGTTCCTAGGGATCA GGACAGTCATCACACCACCGGCAGTCTCCAAACCAAGAGACAGAGGGGTGACATCCAGGA GGAGAAGATCCTGAACCTTCTCATTGCCCTCACCCTTAAGATTGCAGCCTGAACAGC[C/A] GCACCATATGCAACAGCCTCATCGGGATTAATGCTCTTGCAAAGCTCTTTCCATTAAAGAA GTCCTGCAGCAGTTGTTGAACCTTGGGAATTCTGGTAGAACCAACCAAGGACAACATCA TCAACACTTCTTTGTCCATCTTAGCATCCCTCAAACATTTCTCCACCGGCTCCAT
>Gm03_4 0587775	AATAAGTATAAAAAAATACATAAACAGAACAAATCGAATGGAAAAGATAAAATAATTTTGTTA AGAGAGGTTACCGAAGACGGTGTGATGGGGTTCATGGCGACTTGATTCTTGGCCGCATC ACCGATGAGACGCTCGGTGTCAGTGAATCCGACGTAAGACGGCGTGTTCTGTTCCC[T/C] TGGTCTGTTGGCGATGATTTCAACGCGGTGATGTTGCCACACACCGACGCAAGAGTAGGTG GTTCCGAGATCGATTCCGATAGCAGGACCTCTCCTTTTCCGGCCATGGTTTCGGCGTGCG GCGTAGAGAAAAGAAGAGATCTAAGAAAGAGGAAGAAGAAGATGATAACACAGCGATA
>Gm03_4 0600203	CTACGAAAACGCACAATTGGTATTGGCAGGGTAAATTGGTGAAGTTGAACAAAATGAAGCT AAGAAATCACTGATGGAAAGCGGAAATTGGGGAACGATCTGATCCGTGAGCGAAAGGAAG GAAATCAGAGACGAAATTGAGAGTGGTGTGGTTACCTTTGGACGCGGTGGCAGTGGGCG[T /C]GACTTTTCCCTCTGATGAGTGTGTTGTCGTGCTTCTAGGTCTCGCTGCGGNGTTCCC CCACAGTCCCTCCAACACAAATAAGAAGCAGAATAGTGTCTTACTTAGTTGACCATTTTGCCC ATCTCTATTTCCCAAATTTAGATAATAAGGTTTAAATACCATTTTATTCTTTTAAAT
>Gm03_4 0600256	ATGAAGCTAAGAAATCACTGATGGAAAGCGGAAATTGGGGAACGATCTGATCCGTGAGCG AAAGGAAGGAAATCAGAGACGAAATTGAGAGTGGTGTGGTTACCTTTGGACGCGGTGGCA TGGGCGNGACTTTTCCCTCTGATGAGTGTGTTGTCGTGCTTCTAGGTCTCGCTGCGG[ T/G]GTTCCCCCAGAGTCCCTCCAACAAATAAGAAGCAGAATAGTGTCTTACTTAGTTGACC ATTTTGCCCATCTCTATTTCCCAAATTTAGATAATAAGGTTTAAATACCATTTTATTCTTTTAA ATTTGGCTGATTGTTTTTGTATTTAAAAAATATATTTTATTAGTCTTTCAAA
>Gm03_4 0603941	GTCGGGTCGTGGAAAGGGAGGCAAGGGTTTGGGAAAGGGAGGAGCCAAGAGGCACAGGA AGGTGCTTCGTGACAACATCCAGGGTATCACCAAACCCGCAATTCGGCGTTTGGCTCGCA GAGGTGGTGTAAAGAGAATCAGTGGTCTCATCTACGAGGAGACACGTGGCGTCTCAAGA T[C/A]TTTCCTCGAGAACGTTATTCGCGACGCTGTTACCTACACCGAGCACGCTAGAAGAAAG ACCGTTACCGCTATGGATGTTGTTTATGCTCTCAAGAGGCAGGGCAGGACTCTCTACGGTT TCGGTGGTTAATTTCTAGGGTTTCCATCGGCGTAGCAGCTGCTTATTGGTTATTGTTGTAA
>Gm03_4 0628097	ATTTGCAATAGAATGACGAGGAGGATTGAGGACTTCGTGTCCACGTTGTTCCCTTCCCTTG TACTTTGCTGCCAGTGGTTTGAAGAACTGACGTGACTAAGTTACGAAGCGTGGTGGATTGGG GGCTTCTTTTGTGTTACGTCCACCGCAGCGTGGGGAAGATTTTGGGAACGTTTGC[C/G] JGTGGCGATGATGTGCATGGTGTGGGAAAATAAAATAAAATGAAGGAAAATAAATATGAAGA AGATGCACAGTCTTGAATTAATAACAAAATAGCAGTAGCAAATTAATTTAATTGACAACAC ATAATAATGCATTATATCACACAATGAGCACAAGGAAAAATTAATTAAGGGAA
>Gm03_4 0656449	ATTAGATCAACCTTGACCTCACGAACATATGAGCAGGGAAACACAGGATGTCACCTGGAA CATCCAAGGGCATTCTAATGGACCCACCACTGCTGCACTTCTGAGAGAATTTAGAGGAAGG CGTTATGGCGGAGACAAACGCCACCTGCGGCGACATAGACGCCGCTGCCACCATCAAC[A/ G]TTAGCAGCGTTATCCACGTGGCGGCATAGGTTAGTACAGGCCATGAGAGGTGCATACGG ACCATGTTTGAGAACCAGAAAAAAGACATCTTGGGAAGCTCTATGAGGTAGTTGAAATCGT GGGTGTTTCGGGTGGGCAATTTGTCTTTATAAACAAGACAGGAATGCAGTGAATTTGGT
>Gm03_4 0676583	CCTTTCTAAAATTTCAGTATTTTCTCATTGAGCAAAAGCATACATTGCTTACAACTATACCT TCAAACCTTTACTAACAGCCAAGTAATATAAATCAGATAACTTACAGGCAGAGATACAGAGCG TGAGTAAGTTTGATGAGCAGAATGCTCATAGAGTGTTCTGTGACTGAAGTATCA[G/A]CCT CCTTATCAGACGAGCTTGAAGGCAACATGGTATTAGGAATTGGAAAGTTGCAGATTTTTTC AAGCACGCCATATTTACTTTTGTGCTTGATCTTTTATTGCTTTACATATTTTTTCTCATTCA CATTATAGATGTCACCTCCCTCAGAAGCAAGTAAAGATTTCTGAGCATCT
>Gm03_4 0676856	TGATCTTTTATTGCTTTACATATTTTCTCATTACATTATAGATGTCACCTCCCTCAGAA GCAAGTAAAGATTTCTGAGCATCTTGAATGTGTAATGACTCACTAAAAGTCCTTCCAAGATT ACTTATGAAGTTGTCGTCATGCACAACATTGGTATTAGAGCACGTGTCCATCAAT[T/A]GCAT CAGGTACTAAGAGGCAACTAAAAAACAACAATCAGATCAATGAGAATAAATAACACAATTA TCACTAATCTACATAATAAGCATTAAAAATTTAAACATTCCACAGTTATGTTTCATTATTAATA CATTGTGGACCAAAATTTAAAAAAGTTTAAATGATGACAGATG

>Gm03_4 0678154	CTTCATACATTACTAACATAAGTCGGAAGCAAATGATACCTAAGCTTGAAAGTTGAACAAG ATTACCAAAAAAATATACCTTATAAAACGACGTTTGGAGCTGCAACTACAACACTATCAAAA CAGGAGCAGAGAATAGCAACCAACACAGAAAGAAACAACAATAGCAAACCTCGAT[A/C]C CGATCCGCTGACGAATTAATCACAGCTCGCTCAGAGCAGAACCAACAACAACGACGACCAA AAGAGAGCGAGAAAAGAAAGAGAAGGTGGAAGAAAACGAAAGACCTAGTGTTAACTAAGAT TACCACGCCGATCACTTTTTTCAAGATTTTTTCAGACGATGAAAAGAGGATCCACCA
>Gm03_4 0683015	CCACGCTCTTCTTTGTTCTCTTCTTCTTCTCTCGCTTCTCCTTCTCGATATCTTCTTCT CGGCCCTTGGGCTTCTTCTCCGCCGGAGCCTTCTCTGCGGGCTTTTTCTCCGCGGGCTTCTT CTCGGCCCTTTGGGGGAGCCATTGGTTGGTGAAGAGGGGAAAGGAGAGGGTTCGGTTTT[T/C]T ATGATGGGAGAAAAAACTGAATTAGATAGAACGAGTGAAGGAACAGAAGAATGTGTGTGT ATATATAGGAAGAGAGAAAAGCTTTTGATTGGAAGGGGTTGAGTCACGCCGATCGGTGACGT GGCGTCTTGATTGACATTGCGTTTTTTCAGCGCCCAAGGATTGCTTCCGTCTTTT
>Gm03_4 0685721	ATCTACCATCAAAACAGCTAGCCTTAATGAAAAGGCAAAAGTAAAAAATTGCTATAAGATTA AAAAACTGAGCAAAAGATAAATGTATTATTGTAGAGAAATCTAGGTCTCGCATCCTTTACATG TAATTTAAAGCTCAAACAAATTCCTCGATGCGGGCTTCTGAGATTTAGACTCTGAA[T/A]AGC AGGGGAAATCAATCTCTGCTGAACACGGGAAATGAGATGTTGGCTATAACCTCTGCTCGCA ACACTTTGAACCTGAATTTCCGATGAAGAATCATGTCTTATCTAGTACCTGTCAGGCTGTCA CAGCAGGGGAAAACATGAGACAAGTCAAACGGGTAATAAAAAATAAAAAATGT
>Gm03_4 0785291	GGCCTTACCCATGGATTCTTCTCCCAAATTGCGCCACCCTTCTCCATTTCCCACTCCCTTAT CCTCTTCCCTCTCCTTCCGCTCTTTACCGCCACCCCATTTTCTTCGTCTGTCGTTTAGGTTA CCCTCCATCAGAGCCTCCTCCTCACCTCCCAAACGACCCCGCTTTTCGCACCCC[A/G]A GAAACANGCCCTCCCTCCCCCCCCCTCTCCAAACCTAACCTCCTTCTCCCCCTCCT CGAAACCACCTGCATCGTCATCGCCGCCGCCGCTTCTTCTTCATGCGCTTCCACCACACG CCCGCGTCATCGCCGCTCCGCTCACTTCCCCCGCCGCGGAAACCGACACCGCCTT
>Gm03_4 0785299	CCATGGATTCTTCTCCCAAATTGCGCCACCCTTCTCCATTTCCCACTCCCTTATCCTCTTCC CTCTCCTTCCGCTCTTTACCGCCACCCCATTTTCTTCGTCTGTCGTTTAGGTTACCTTCCAT CAGAGCCTCCTCCTCACCTCCCAAACGACCCCGCTTTTCGCACCCCNAGAAACA[A/C]G CCCTCCCTCCCCCCCCCTCTCCAAACCTAACCTCCTTCTCTCCCCCTCCTCGAAACCA CCTGCATCGTCATCGCCGCCGCCGCTTCTTCTTCATGCGCTTCCACCACACGCCCGCCG TCATCGCCGCTCCGCTCACTTCCCCCGCCGCGGAAACCGACACCGCCTTACCCAGG
>Gm03_4 0823593	TACATCTTAATTCGAAAATCTATCATCGACGAAGGAATCTACCGCCTCGGCGTGAGAAAAC TTAGCTCCTCGCGCGCTAACAAGATGGATTGGGAAGTTCTTGATTTGAAGATCAAGAGTTG GTTAGAGGCGACAAGGATTTCCGTTAGAACACTCTTTAACGGAGAGAGAATACTCTGC[G/C] ACCACGTCTTCAGTTACTCAGATTCCGTCAGAGAATCATGCTTCGCCGAGATTTCAAGAGA CGGTGCCGCTCTCCTTTTCGAGATTCCTGAACTCGTTGCCAAAACAAAGAAATCTTCGCCG GAGAAGCTGTTCCGCGTGCTCGATATGCATGCTGTGGCTTCCGAAGTGTGCCGGAG
>Gm03_4 0874888	CCTGGGGTGACCGTACAATATCTGAGAACCCTTTTCTCTGTTGCAACTTCTAGACCTCACGA ACTCAGCATCCGATTCCGGCCACTTATAGAGGTTACGTAGGAAGCCCTCACCGGAACAC GTGGGTCACGTGCGTCGTTGATGCAGTTTATGATATGCATACTGAGCTCATCTTTTTTTT[C/T]C CCTCAGTTTCTCAGGTGAAACTGTGATAGCTAGTGGGAGTTTGTGTAGAAGGGTATAAATT GAGAATATTAATAAAAAAGAAAAAAGAAAAAAGAAAAAGGAGAAAAAAGCTTAGATTGCAAGAAA AGGGTGTGGGAAAGGAAAGAAGGACTAATTGAATGTATTTATAGTGGGCATTATT
>Gm03_4 0886333	TGTGGAATTAATCAGTTCCACTTTGCGATTGGAGCAGGGTTCTTTGACTGGAGAGAGTGAA GCTGTGAATAAGACTAACAAGCGTGTGGATGAGGATGCTGATATTCAGGGGAAGAGGTGT ATGGTTTTTCGACGGGACCACTGTTGTTAATGGGAATAGCATATGCTTGGTCACTCAGAC[T/C] JGGCATGGATACCGAAATCGGGAAGGTGCATATGCAGATACACGTAGCCTCGCAGAGCGA GGAGGATACCCCATTTGAAGAAGAACTCAATGAGTTTGGAGAGAAGCTGACCTTGATAATT GGACTGATATGTATTTGGTTTGGCTCATCAATGTTAAGTACTTCTTTCTTGGGAGTA
>Gm03_4 0889026	CTCCTAGGGAGGAAGTATACCAAGCAATTGAAGACTGCAGAGATGCTGGAATTCGTGTTAT GGTTATTACTGGAGACAACAAGAACACCGCTGAAGCTATATGCCGTGAAATAGGTGTATTTT CTCCAGATGAAGACATTAGTTCTAAAAGTTTAAACAGGTAGAGATTTTATGGAAGTGC[A/G]TG ATAAAAAGGCATATCTGAGACAGCATGGCGGGCTTTTGTTCCTAAGGGCTGAACCTAGGCA CAAGCAAGAAATTGTGAGGCTGCTCAAAGAGGAAGGGGAAGTGGTGGCCATGACTGGGGA TGGTGTTAATGATGCTCCTGCCTTGAAGCTTGACAGACATTGGTATTGCAATGGGCA
>Gm03_4 0906651	TTAACAGTCACTAATCTTATTGGTTCTGATGCAACGACTCTGCAATTTCAATTGGCTGTAGTT TCAGTAGTCAGAATTGTTTTATAACATCTTTGCTTGTAAGCTGAGTTTTGGTTACCTTGGTTC TTGCCAGCCGATCGAGATCTCCTGTCAGAAGAGAACGAAGCCAGTTTCTGAAG[A/C]TAGA

	TCACAGAGTCGTGAGCCCTCCAAGATAAGGAAGCATAGTGCATCACCTGACCAAAGCAGC CCACGGAAGAGAGGTGACACATCTCCTGGCAATGATAGGTTGGCCACTCATCAAGATGGG TCTGACTACAGTGATGGTCCTAGAGGGAAAAGTAGAAGCCCTGCTAGCCCTGCAA
>Gm03_4 1007691	CAAAAAAAGAAGCCCACTTAGAAGGACTCTTTAAATATATTTATATTCAACGTATAAAACCTT TGATGCTGTTCTGCCAAACCCCAAACCCTAGTTTTGTCAGTAGCACACTTCACCCATGGCA GTAGCAGCAGCTCGCCCTATCGTCTCCGTCCAAACCCTAGAGGGCGATGCAACCCC[T/C]A CCGTTCTCTTCCCGACGTCATGAAAGCCTCCATTGGCCGTGACATCGTCAATTTCTGTTCA CTCCAACATTTCTCGCAACAGCCGCCAACCCCTACGCCGTGAGCAAGCGCGCCGGCCACCA AACCTCCGCCGAGTCCTGGGGCACCAGGCCGCGCCGTCTCGCGTATCCCGCGAGTCCC
>Gm03_4 1007937	CAACATTTCTCGCAACAGCCGCCAACCCCTACGCCGTGAGCAAGCGCGCCGCCACCAAAC CTCCGCCGAGTCCTGGGGCACCAGGCCGCGCCGTCTCGCGTATCCCGCGAGTCCCCGGCG GTGGTACCCACCGCGCCGCCAGGGCACCTTCGGAACATGTGCCGTGGCGGCCGCATG TT[T/C]GCTCCCACCAAGATCTGGCGCCGTGGCACCGCAAGATCAACGTTCAACAGAAGC GCCACGCCGTCTGTCGCCCATCGCCGCTCCGCAATCCCTCGTCTCGTCCAGGCTCGCG GCCACCGCATCGAGTCCGTCCCGGAACTCCCCCTCGTCTGTCAGCGACACTGTGAAAGCG TCGA
>Gm03_4 1008255	GGAACCTCCCCCTCGTCGTGAGCGACACTGTGAAAGCGTCGAGAAATCCAAAGAAGCCGT CAAAATTCTCCAGAAAATCGGCGCATTACAGACGCGGAGAAAGCTAACTCAGCCACGGA ATTCGTCCCGGGAAGGGAAAAATGAGGAACCGCAGGTATATCTCTGGCAAGGGTCTCT[T/ C]ATAGTGTACGGAACAGAAGGTGCTAAGGCCATTAAAGCCTTCAGGAACATTCCCGGTGT AGAGGTGGCAAACGTTGATAGGCTCAACCTTCTGAAGCTGGCACCCGGTGGCCACCTTGG AAGGTTCAATTATTTGGACCAAGTCCGCATTTGAGAAATTGGACTCTATCTATGGATCCTT
>Gm03_4 1008442	TACGGAACAGAAGGTGCTAAGGCCATTAAAGCCTTCAGGAACATTCCCGGTGTAGAGGTG GCAAACGTTGATAGGCTCAACCTTCTGAAGCTGGCACCCGGTGGCCACCTTGGAAAGGTTT ATTATTTGGACCAAGTCCGCATTTGAGAAATTGGACTCTATCTATGGATCCTTCGATAAG[G/ A]CCTCCGAGAAGAAAAANGGCTACTTGCTTCCCAGGCCCAAGATGATCAACTCTGACTTG GCCCGTATCATTAACCTCCGACGAGGTCCAGTCCGTGGTCAGGCCCATTAAGAAAGATGTTA AGAGAGCACCTCTCAAGAAGAACCCTCTCACGAACCTCAATGCGATGTTGAAGCTCAAT
>Gm03_4 1008459	TAAGGCCATTAAAGCCTTCAGGAACATTCCCGGTGTAGAGGTGGCAAACGTTGATAGGCTC AACCTTCTGAAGCTGGCACCCGGTGGCCACCTTGGAAAGGTTTATTATTTGGACCAAGTCCG CATTTGAGAAATTGGACTCTATCTATGGATCCTTCGATAAGNCCTCCGAGAAGAAAAA[T/G] GGCTACTTGCTTCCCAGGCCCAAGATGATCAACTCTGACTTGCCCCGTATCATTAACCTCCG ACGAGGTCCAGTCCGTGGTCAGGCCCATTAAGAAAGATGTTAAGAGAGCACCTCTCAAGAA GAACCCTCTCACGAACCTCAATGCGATGTTGAAGCTCAATCCCTATGCTAAGACCGC
>Gm03_4 1171468	TTCCGCAAAGAACCCTCACCTTTATTGAAAGGCCAGTACCTCTCAATGTGGGTGTCATAG TACAATAATGATCCCAGGGAATATATTGACAATCAATATTTGGCTTCTCCTGCCACAATCT GAAGATATAGGAAGATCAGTTGAACTCCAGCATTAAGATATCTACTAGTACTGCCT[C/G]C CTCATAGACTGAAGGAATTCCAGGGCTCTTACGCTTGATGCCCTTGTTCACTTCCCAT GTAAGATCAATAAAACCACCATCAACATGAAATGTTTGTCTGTCATAATTGGAGGAAGAAGG TTGGTGTGCATAATTTTCAACGGTATAAACAATGATGGTCCAGCACTAGACAAA
>Gm03_4 1185505	CTGGCATGCAAATGTTCAAAATTAATGGGGTTATACCTTTGTCCAATGCATGGTACAATGTA TTATTGCAGGTAAAGGGGCTTTCATCACAGAATTGCTGCCCGTAATGATTTGGCTCTTGC ATTGCCGGATAGGATTCAAGCTATCCAGATGGGACCCCTGCAGCATCCAAACAAA[G/C]T GGAAGTTGGGCAGCTTCAGCCTCACGTCTGGAATTAATTTGATTCAGGTAAATTGAAATTA TCCATGTCTCATAGAAGTTTATCAATGGGAGACTTTTTAATTGATGGTCATCCATTCATTTT GTGTATCTTGTCAATTCCAGATGGGAAATTCGATGATGAATACTTCCAACAAA
>Gm03_4 1223681	TATCAGCTATTTGTGAGGCTTTGAGGAGTTTGATGAAGGTTGTGAATTGCTGCTTTCTCA AGCACTTGGGTGATGTTGGTGGGTCCCGCTGGTGCCGGTGCTACTTGGGCTGAAATTGTTT GGATGAAGATGTTGAGGAAGAGGAAAGGGAGGAAATTGAATAGCTTTGTCAATTGGTG[T/A]T CCTTGTGAGTGAGGAAATTAAGAAGAGAAGACTAAGATGTTTGTACAAGTGAAGGAAGA GATGTGGTGTGTAATATATATTAAGAAGAGGGTGCATGGAAATTTGATTGTTTAAAGGTT AGGTGAAAGGTTGCAATTAGTATAAGGTAGCAAGAGGCCCTCATGCAATCAATTT
>Gm03_4 1228284	TAGGCCTTGGTTTGAGTTGTTGAGCTGTGAATTGATTCTATCAGCTATTTGTGAAGCTTTGA GGAGTTTAATGAAGGTTGTGAATTGTCCTGCTTTCTCTAGAACTTGGGTGATGTTGGTGGG GCCCCTGGTGTGCTGCTGGTGTGTTGGGCTGAAATTGTTGGATGAAAAGGAAAAGGAG[A/G] AAATGGATTAGCTTTGCCATTTGTGTGAGTGAGGAANCTAAGAAGACTAAGATTTTTGTCAC AAGTGAAGAAGGGGATGTGGTGTGTTGTTATATTATATGGTGTAAAGGTTTGAGGGT

	GAGTGGAATAGATTGGTATAAAAAAGTTAGGTGAAAGGTTGCAATTTGTATAAGG
>Gm03_4 1228321	CTATCAGCTATTTGTGAAGCTTTGAGGAGTTAATGAAGGTTGTGAATTGTCCTGCTTTCTC TAGAAGCTGGGTGATGTTGGTGGGGCCCGCTGGTGCTGCTGGTGTGGGCTGAAATTGTT TGGATGAAAAGGAAAAGGAGNAAATGGATTAGCTTTGCCATTTGTGTGAGTGAGGAA[G/A]C TAAGAAGACTAAGATTTTTGTACAAGTGGAAGAAGGGGATGTGGTGTGTATTATATT ATATGGTGTAAAGGTTTGAGGGTGAGTGGAATAGATTGGTATAAAAAAGTTAGGTGAAAG GTTGCAATTTGTATAAGGTGGCAAGAGGCCCTCATGCAATCAATTTTTCTTTCTA
>Gm03_4 1234338	GATGATTTAGTTCAAGTTTGGAGCATGGAAGATCGCAAGGTTGTGGCATGGGGCGAGGGA CATAACTCATGGGTAAAGTTGCCTGACAATTTTTAGTTGTGTGCTTGTAGCAAACCTTTGCTAC CGTAGGGTTAATGTAGTGTGTATCAAATGCCATGATCAGGTGAGTGAGTTGCTTT[T/C] GATTCATATTGGACATCACCAAAATCAAATGACAATGGAGAGACTGTTACATATCGTTTTGG TTCTGTTGGTCAGGTGCTCAGAAAAATGTTTCATTATAATTTTTACTTTTTCTTTGTGTAGCTT TTCATTCTAAGTAAAACCTATTGGTATACCTAACATGGATCTGTGGTAGCTTT
>Gm03_4 1274006	AATATACACTGACAAAGTGATCCTCAGACCTCTATTGAAACGGTGATACATATAGGCATGCC TATGAATCTTAACTGTTTATTTTTAAAACTCAGAAATATTTTTAATACATGGCAGCAAGGT ACCACGTATTATAGCCAACAAAAAGCAAGAATCTATCAGCGCCAGGATCAGA[A/G]TCAA CAATAAACACAATATAAATGAGATATACTGTTCTGCATAACTTAAGAGTTCCCCAGCACCAA TATTTCTCTAATTCTTCACACTTGTTACATGCTTGAGGAGCATGGAGACAACATCCCCTGAT ATCCTCTCAGCCTCAGTTTTCAGGTGATGAATCTTTCTTCTGTCTCTTGC
>Gm03_4 1275788	GAAAAGGAAATCGTGAGAAACGGGAGCGAAAACGAGTTGATTATAGTAAGAAAGCGCGTG AAATCCGAAAAGGGCTGCATTGGCGTGAAAATGGTAGATCTAAAAGTAAAGGAGAAAACCG AAATAAAGAGATTGGAAGGTACCGTAGCAATCGGGTCTGTGTTAGATTCAGAGAAAACA[A/ G]GACACAGTGTCTCTAACGAAAACCTGGAGGACTGCTTCGCTGATTTGCTTTTCTGCTTCCG CCAATCAAATCAACACAACCCACCTATTTCTTTAATAATAAATAAATCACACATATATAATTTA ATTTTACTCACAAACCATATTCTTTCGCAAAGTGTGGACATAAATAAATGACA
>Gm03_4 1982791	TGGATATGAGTGGAACAAATATAATCAGACTCACTATGATCATGACAACCCACCTCCAAAGA TAGTCCAAGGGTACAAATTTAATATTTTCTACCCTGATCTTGTAGACAAGACAAAAGCCCCA ACCTACACCATTGAGAAGGATGGCAGCAATGGGAGACTTGCAATTATAAGATTCCA[C/T]GC AGGGCCACCATATGAAGACATAGTAAGTCTATTATCCTTTAATTTTATGAAGGATTTTAAAT TTCAATCAATTTACTTATTTCTTATTTTTCCATGACAGGCTTTCCGCATTGTTAACAAAGAA TGGGAGTATTCTCATAAGAAAGGTTTAAAGTGTGCATTGAACGTGGAAT
>Gm03_4 2142835	CCTTGTTATGAGAACTTAAGGCATCTTCTGTCTTAAATGACCTGCAAGCCAATATAAAAAATTA TTGTGAATTGATGTGCCTGATTACTAAGAAAAAGGAAAAAAGGAAAAGGAGGAGACAAAT CAATATAAAAAGACATCTCCAATCACCTGTTGCAAGGCTTGACAGCTGTAATCTCCA[C/A]CTG ACTTCGGAGTCTGCTGCTTGTATTAGCAGTTGCTTTACCAGTCTGCTTAGAAGGGTGAGG AGTTGCAACATGGACAACATTCTTGCTATCTGAACAGATCATTAAATCTCATCAATATAGTTT CAGTTTATTGTGGTCACAACATGCATGCACAGACAAGTAAAACAATCAAACA
>Gm03_4 2179147	CTCCGCGACCCTACCCTATCAATTTTCGTACTTCTTTTCGTAATTGCAGCGCAAGAATGGCGA AGAGGAAGGAGCGGATTGAGAACCCGAGCCATTGATCCGTCAGCGCGCAGCCCGCGC AAGACGAAGAAGCGTTTCGAAGCCGCCGAAGCGGCACCAGCAGGAAGAGCAGTTTCATTGC[ T/C]CCCAAACCTCAGCTCCAAGATCATGAAGCAGGCGCTGATCCAGCAGAAGGAGGAAGAG AAGGAGGAAACGCACGAGAACAACGCCGCCGCGAATTTATTCGAAGAGGTTCCGAACCTC GAGGAAGACGGCGGTGACGACATCGACGACTTCGCCGGGTTCTCCGAAACGCAGAGCCA GTT
>Gm03_4 2224757	CGTATAAGAATCTCGAATCCGCAAGTTTTAGATCCATTGAAATGAGCAATAGTAACAGTGCC AACGAATGACGCAAGGGGTGACGCATGTGCTCTTAGTTGACAATTGGTGGGGTCCATCTAA TCTAATGTTTGTCTTCTGTTGGTGTGAGGTCCGGTGAAAAGGTACCGGATTGTGTT[G/T]GT TTTCTCACCACCCAGGGGNGCAACTCTCGCAAGTGCGGTTTTCAAAAACATGACTTCACTA CAGTGTTATGATCATAGCATGGTAAGTCAATCAGCTTCATAGACTGAACCTAAACTAAATA AACATTTGTCGGACGTGTCTTTTTATTTTCAATTTGACTTGAAACTGCCGCC
>Gm03_4 2224778	CAAGTTTTAGATCCATTGAAATGAGCAATAGTAACAGTGCCAACGAATGACGCAAGGGGTG ACGCATGTGCTCTTAGTTGACAATTGGTGGGGTCCATCTAATCTAATGTTTGTCTGTTG GTGTTGAGGTCCGGTGAAAAGGTACCGGATTGTGTTNGTTTTCTCACCACCCAGGGG[C/T] GCAACTCTCGCAAGTGCGGTTTTCAAAAACATGACTTCACTACAGTGTTATGATCATAGCAT GGTAAAGTCAATCAGCTTCATAGACTGAACCTAAACTAAATAAACATTTGTGCGACGTGTCT TTTTTATATTTCAATTTGACTTGAAACTGCCGCCGCCGCCGCCGCCGCCGCCGCCGCC
>Gm03_4	GCTCAAGATCGGCCCAAACGAACCATCCGAGCTGGCTATCCATGAAAATGCCTACGGTCTT

2243466	GCCCGATATGCCGTCATATGCCAGGAGAATGGTCTGGTACCTATTGTAGAGCCAGAGATCC TGGTGGATGGACCTCATGACATCAACAAGTGTGCTGAGGTGACCGAGCGCTTCTTGC[A/T ]GCATGCTACAAGGCTCTAAATGATCACCATGTTCTGCTTGAGGGCACTCTGTTGAAGCCCA ACATGGTCACCCCTGGTTCAGAGTCTAAGAAGGTACCCCCAGAGGTGATTGCTCAATACAC TGTTACAGCTTTGCAGCGAACTGTTCCCGCTGCTGTTCCGGCCATTGTCTTCTTGTG
>Gm03_4 2243699	TGAAGCCCAACATGGTCACCCCTGGTTCAGAGTCTAAGAAGGTACCCCCAGAGGTGATTG CTCAATACACTGTTACAGCTTTGCAGCGAACTGTTCCCGCTGCTGTTCCGGCCATTGTCTTC TTGTCTGGTGGGCAGAGCGAGGAGGAGGCAACCCCTCAACCTCAACGCCATGAACAAGT[C/ A]CCAGGGAAAGAAGCCGTGGTCCCTTTCTTTCTTTTGAAGGGCACTTCAGCAAAGCAC TCTCAAGGCATGGGGTGGGAAAGATGAAAAACATTAAGAAGGCTCAGGATGCTTTATTTGCC AGGTGCAATGCAAACTCACATGCAACTTTGGGAACCTACAAAGGTGATGCTACCCTTG
>Gm03_4 2672131	CCCCAAAACAAAAAAGACAATTGCAGATCAATACGCAGCGAATGTAGAAGAACATGGAAA ATTAATAAATAAACTAAAAATTAATAAATTAAGTGGAAAGTGAGAGAAACACTAACTTCT TCAAATCCGGTGGTTGCCCTGATCTGCTCATTGTGAGGAGAAGAGATCACAGATGC[C/G]AA CGAAGGAAGAAAAATTTGGTTTCGAAGTGGCAAAACCTCAACTACGCTCACTCAAGACCCTA ACATCGGAGGCTATACTCTTTATCGTTTAAAGCCAGGCCAGTATACGAAACGATTGCTCAA AGGCACATGGATCTGATAGCATTAGTAAACAGCCAGGACCAAAATATGGGAAAG
>Gm03_4 5026222	CCTTTGGCCCTTCATCATAAAAAGTGTGTAAATTGCTTCCTCCTCTTTGCTTGTGTGGATT TGAAAATGTTGAAAACCTGAGAAAAAAGAAGATTTCTTTGCCTGTTTTTGCACCCCATGGTT ATGATTGTTGATTTCTTGGCTATGAAAGACTATAGTGTCACTTTTAGTGACTAGA[T/A]ATAG GAACCTGTAGTATTTGTGGTGTGGGTTAGAACCTCGAGGCTATGGGCCTCTCAATTTATAG TACAAGTTAGGCCACTTCTTTAACTGAGATTAAGGGTGGAGAATCTGATTCAAACCAGTGT TTATGGTCCAATTAAGCTGACATGGAACATTTGAAACTGTTCCATTTCAT
>Gm03_4 5416367	GGTTGCTTGTATAGATGTTGATGCTGCCCTACCGTTCAATATTCCATTATCCATAGGTGGA CTGGTTCCCACTTACCTTCTGATTCACTAGGCTCTATCACTTTCACGGACCCATCTGTCAA TCCAACAGCAAATTGGTTGGGTTCCAGTGGATGAGCCGCAATGGCAACAGGATACG[T/C]A GATTGATTCTGCAATATAGTATACCACAATATGGATCACCAAATGGAAGTAAGACTTCACG ATACTACATCCTTAATAAATGCTCAATGTTTAATACAATAGAGAGTAAATTTGGGTCTGATC TGTTGGGCAGAACTTACCCGCTTAAAGCTGCTGGAGGTGAAAAGTATGTGGA
>Gm03_4 5503654	CTAAATACACAAATAGAAAATGTTTTGAAAAGGAGCAGGGAGTGCAACATTATAAAGAGTG GCCACTCAATAGTCAATAAGGGGAATCAAAAAATCAAATGCTAGTTTTTGTAAATACACGTC AATGTAAACATGGGAGCCACGATATGTTATCAGAACAGAAATATCATAAAAACTGT[T/C]AC CGACGCCTAGAGCAAACAGTTTAATTTCAAGAAGTGAACAATTGTAAAAAATAAGAACAA AGTTATATTATTTCTTAACTCTAAATCCTCTGCTGTTCTCCTCCTCTAGCCCTCTCGA ACTTTCTTCCCTTTGCTCGTACATAGGGCTTGGTATGGCTGTGTGGCACACC
>Gm03_4 5516926	AGGAGGAATGAACAATACCAAAAAACAACGAGAATGGCTTATTAATTATAAATATACATTA CATGATAAGGAGTAAAGATACACCGCTACTCTACCCTTACACTTTCTACTTACGATTTGGCA ACCACTACGCAGCTTACTTAAGTAAAGACTTAAAGAAGCTAAACCAAAGCTACCT[T/C]GC ATGAACCTGGCAGAACTTCTTGCAGACATTGAGACATTATACATGCACACAAAGACTAAAC AAAAATGAATACAGCAAAGGCCAAAAAAGCTGCTACATCCTCAGCTTTGAGGTACGAGAAA ATGCGTCACACATCGAAAGGAGAAGAAGGGCCTGATCTGTTCTGCCCTCTCTTC
>Gm04_4 4787569	GGTGGGGATAGGCGAAGGAGGTGGCGACTGAGTAGTGAATAGGGTTTTTGATTGATTGAT AGCAATAATTGGTGTTAGCGAAGCTTGAGAATTGAGATTTATGGGTCGCAATCGCAGTCGC AGTCGCAGTGTGTCACGCTCTCCTTCCAATTCCCGCAGAAAGAAGGTACTCTCCATCCCC[T/ C]GTTTCTCATCGCCACTCGAGAACTTCCAGGCGCCGCTCTCCTTCCACAAGCGCCGTCG AAGGCACAGAACCTCTTCTCCCCCTCTCTTTCACGCAGCCCCACTCCCAACCCAAGAAA GATCAAAACAAAAGGTAACCAATCATAACCATTCCTTTATTTGCCTGGATTTTATCTTA
>Gm04_4 5061509	AGAAAGAGCTGTTCAACTTTGTGATGAGTCACTTGGATGCAGCCACACAAGATGAAGAAGG GTCTGCACCTGTTTTGTTTCAACCATTTGGGAGCTACCTTTTTGTTTCATCAGATGGAGCAG TGTGTGTGGATTGTGCAACTTCTGTTATAAAGATGTTGCATTTGATGTTTGCATCAG[T/G]TT CCCCAGCTTGTAGTATTGAGGACCATCTAAAGTATGGAGACTATGTTAAGAATTTGTCATTA CAATTCCTGAACCAGAACAAATTCGTGCAGGGGAATATTCCTGATTCAAGCTATGAAGCAG GACTTGAATTGTCGGTTCAATCTTCAGGTTTGGGTAACCAGGTAAATGCACTAT
>Gm04_4 5090419	CCAACCCCCCTCCCCCAATAAAAAATAGGAAAAGAAATTAGCTAACCACTAATGACTGATGAT GTAAACAGTTATGGAGGAACCGTTGGCAGCTTGAAGAACATTGCGACTAGAAACATGTCAA TTGCTTTGCAAACCAAGGCATACAATCTGACTGTTTTTGCCTGTATTCTTACAAATT[C/A]TG TGCAAAATTGGTGCAGTTACTGGAGTTTGATTGGTTTTACTATTTGCAGAGACATTGAATTT

	GTACAGATTTGCATGTATAATAATACATCAGTTTTAACAAGAATTTGTAAGTGAAGAAATCACA CGCCCCGTCCCTCTTGCAACTACTTTCACATTATTTACGTTAGATACTTCC
>Gm04_4 5152573	ATATTGGTTTCAGAATTAGCTGAAGTACTATCAAGTTTGTCAATATCATGCCAAACACCAGT CCTTCTTATCTCCATCTATCAAAGGTAAAAGTTACCATAAAAGCAGACAACATTGAATTTTCA AGTGCTATCAAGTACCTGAAAAGTTGTCGAATAAGCGCTCAGTAATGTTATCATCA[T/A]AGTA CTTCATCTCATCCCATGGTCCATCTCCAACACCAACCAAAATAATCGAGAGAGGATAATGAC TGATAATCAGAATAGCTCCACATTTAAGACAGATACATAAAACAACAAAAGTTACAAAGATA TCACCAAAAAAATAACAAGAGAAGTTACTCTACCTGGCAGCAATGATAGAA
>Gm04_4 5157974	GTGGATTTGTTTAGCGTGGAGAGGAAAGCTTTGTTGCAGAGTTTGCGTTTGACGATGTTTG GGTGTGTTGAGATTGGCGAACAGAAAGATGATGATGGCCATTTGTTAGACTTGTGTTGCGT TGCGTGCAC TAGTTGACTACTTCCCTTTTACATTCCCTCTTCCCTCGTGGTAGCACCA[T/A]A GGTGCATTCCCTATCCATCCCTTAGTCAAAGCTTTTCTAACTCTAATAATTCCCTTTTACTAAA ACTTTCCCTCTAGACGCTTGCCACCTCATTCTCAGGGATCCTTCCCTCTGCCATTTTTTTTTT TTACTAAAATGGGTAAAATAAATAATTATAATAATACGTTGAACAAA
>Gm04_4 5380790	GTTGTTTTCTTTCACAGCCAATGTGTCATCATCTTCAAAGAAACAATCAAGCAAAGATAGAA TGTCATTGTCAACTTGATTATTTGGAGCTGAAGGATGTGCTGAGCAAGAGGTCTCAATTCT GGTACACAAGATACAGAGCCAATTGATTGCTTGGTGTGAAGGATGGTCTGAGAA[T/A]GA GGGCATCAATTCTGATGCAGATGACACAGAGCCAATTGACAGTTTGGTGTGAAGGATGT GCCGAGCAAGATGGCATGAATTCTGAAACACGTGAAACAGAGCCAATTGACTGTTTGGTG CTGAAGGATGTGCCGAGCAAGAGGGCATGAATTCTGAAACACATGAAACAGAGCC
>Gm04_4 5594453	AGGTAATAAAAAAAGTTAAACAGGAAACTTTTTCGCTTTTTCAATAATTGGTTGGAGACAAG AACTATTTTTGTGGAAGACATCTTAGACATGTATAGTTTAGTTATTTCAATTTTTCTTAAC AGGAAACAATCATTAGTAATTACTCTCATAAAATTGGTGATGCCAGCAACCAA[A/C]TTTTG GTGTTGACTGCGATGAGCAGTAGTACATTAGGCCTCATTGAGGTTTATGACAATCATGGCC GTGCTTACCCTTGCAAAGGAAATGAACATGAGAAATCTTAATCTTATAGATCCCCTAGAAAA CCTTAATAAACTCATCTACAAATCATAGAAATGGACTCATAAATTCTGTAA
>Gm05_3 8251772	TGCTAATATTTCTTTGATACCCTCTGTAACCCCAAGATCTGTCTGGTAAGGTTGATTGTGGC CAAATGAAGATGGTAAACTTGAGCAGCTTCACTCAGATGAAGCATATGAGATGATTCAAAAC CACTTGAGTCTCATTCTCGGTAACAAAGCTGGAGATTTAACATCAGTAGCTGAAAT[C/T]AGC AAATTCAGGGTAGGGCAGGTCTATGCAGCATCAGTGATGTATGGTTATTTCTTAAGCGAG TTGACCAAAGGTTCCAGTTGGAGAAGACAATGAAAGTTCTTCCAAATGCAACAGAAGAAGA GAATGGTGTTTCATCGAAATACGATGGACAATGCAAGACCCAGTATTGAACAGGA
>Gm05_3 8278658	AGGTTCCCTCCCTCAGATCTGGCCTCCTGCGTCAACCTCCGCAACCTCTACATTCAGCGC AACTTGCTCACCGGCCAAATTCACCCTTTTGTTTCATTTGCCGGACCTCGTTCGCTTGAA CATGGGCTTCAACAACCTTCTCCGGCCCCCTTTCCGTCCGCTTCAACAACCTTACCCG[T/C]T TGAAAACCTTTGTTTCTCGAAAACAACAGCTCTCCGGCCCAATCCCCGACTTGAACAACTC ACCCTCGACCAAGTTCAACGTCTCCGACAACCTCCTCAACGGCTCTGTCCCTTTGAAGCTTC AGACATTCCCTCAGGACTCTTTTCTAGGTAACCTCCCTCTGTGGCCGACCACTCTC
>Gm05_3 8279366	GGAAGGGAACGCGAAGAAGTTGGTGTGTTTTGGGAATGCGGCGAGGGCGTTTGATTGGA AGATTTGCTCAGGGCTTCGGCGGAGGTTTTGGGAAAGGGACTTTCGGGACGGCGTACAA GGCGGTGTTGGAGGCGGGGCCGTTGGTGCCGTGAAGAGGTTGAAGGATGTGACAATTT C[C/G]GAGAAGGAATTTAAGGAGAAGATTGAGGCGGTGGGAGCGATGGATCACGAGAGTTT GGTGCTCTCAGGGCTTACTATTTACGAGGGATGAGAAGCTCCTTGTCTATGATTATATG CCCATGGGAAGCTTGTCTGCCCTTTTACATGGTCCGTCTTCTTTCTTTTTTGTGCTTGCT
>Gm05_3 8280387	GGAGTGGACCTCCCCAGATGGGTTCAATCCGTGGTTAGAGAAGAGTGGACTTCTGAGGTC TTTGATCTTGAGCTCCTTAGGTATCAAAATGTAGAAGAGGAGATGGTTCAAGTTGTTGCACT TGCAGTTGATTGTGCAGCACAATACCTTGACAAGCGCCCTTCAATGTCTGAAGTGGTA[A/C] GGAGCATACAAGAGTTGCGAAGGTCTAGCTTGAAAGAGGACCAGGACCAATCCAACATG ACAATGATATACTATTATAGAAGAACATGTATCTTCTCCCTCGACTTTGCTTTGTGAGAGTCG GAGCCCATATTTTACTTGCTAAATCCTGTGCGTATGTTGCTTGCATTCTGTTGGCT
>Gm05_3 8337295	TTTTGTCCAATACTGTTAGTTTTCTAACGAGGGGGAGGAAAATAAAATGAGGGGGCCAAGA GTTTTCTTCTCAAATTCAAATGAGGGGAACAATTGTTATGGCAAATTTACCAAATATTCAA CCAAAAACCAAGACGTTGAAATTTATTTTTAGCAAGGAAGCCAATGGCAAGTGACC[A/T]TT GTTATTCAAATCGAAGAGCATTCTCCTCTCAGCAGATCGTCATTCTAATTCGAAAGCAGCTA TGAATCTTAAACCAACATAAAAGACACAGATAACTATTCCGGCCACACAGGAATTAATATA GCTAGAACAAGGCATATGATAGCAATAAAGCATGTCAATTTTTCTTGGCAATG
>Gm05_3	CATAGGAAAGTACATAGAACAAGTCTACAAGGACCTTCCACCCACTCACCCCTGCTTTGTTG

8402327	ACTCACCATCTCAACCGATTGAAATCCTCTGCCCTCCTCGTCCCTCGTCAAGAAATCCTACAA GCTTCCTGGATCTCATCATCTGCCTGCGCTCCAGCCCCAAAAGACCCGTTGGTCGTCC[T/C] CCCAAGCCCAAGCCCCACCCCGTCTGGCCTGCTGCAGCCCTACCGACAGCCCTGCTCTT CTATCCGCCAAAAGAGGCCCGGTCGTCTAAAAAGATTGGTGTAGCCCGCCAGGCCCA AGGGGTCTGCCACCCGGGACTGGGAGATCTAAGCTGCCAAGAGGCCTGGCCGTCCCC
>Gm05_3 8432746	CATTAGTAAGACTTTCCGTGCGCTGCTTGAGGTAAGTGAGAAACAAACATTTACTGGTCCAC CGGAAAATGTCAGAGATCATGTTATGGCTGCCACAAGGGTCTTAACAAGGGAGACTTCCA GAAGGCTTTTGACATTATTGTGTCTCTTGATGTGTGGAAATTTGTGAGAAATCGAGA[C/T]AC CGTACTTTGAAATGCTGAAGGACAAAATCAAGGAGGAGGCATTGAGGACATACCTCTTTACC TTCTCTTCATCTTATGAATCGTTGAGCCTGGATCAACTCACAAAATTTTTGATCTCTCTGTT TGTCGCACCTATAGCAATTGTCAGTAGGATGATGATCAATGAGGAGCTTCATGC
>Gm05_3 8433427	TGGGAGTGGCTATTCAAGGGGACGAGAAAGGGGTTTCGTATGGAGTTTCAGGGAGGACTG CTCAAAGGGGTTACGATTGAGGGGACCCAGGGAGATGGATCTACTCGCATGGTTAGCC TTAAGGGAGTTCGTGCCTGAATAAGAGTTTGATAATTTGGCTAACGCACTATAGCATTCT[A /G]GTTTTCACTTCTGTAGGCTTTAATTTCTTACGGATTTATCTTTGTTATTTGTTACATTG GAGCTTGCTTTTCATTTTACACCTATTTTACCTGGTTTTATTTGTTTGAATTGTACCTTAAG TGATGCATATTGACCTTGCTCATTTANGTGAGCTATCAATGTCTTTACACAA
>Gm05_3 8433580	AATTTGGCTAACGCACTATAGCATTCTNGTTTTCACTTCTGTAGGCTTTAATTTTCTTACGGA TTTTATCTTTGTTTATTTGTTACATTGGAGCTTGCTTTCACTTTACACCTATTTTACCTGGTT TTATTTTGTTTGAATTGTACCCTTAAGTGATGCATATTGACCTTGCTCATTTA[C/A]GTGAGCT ATCAATGTCTTTACACAAAACCTATTCTTTGTTTTCTTTTGTAAATGAAAGTAATAATTTAAAG AAATTTTTCAATAATTTTCTAATTACCTAATTTATCATGTTTATAAATATTAGGCTTAAATATTA TTTTAAACAATAATGTAATTTTGGGCTCATTGTGTAGAAC
>Gm05_3 8589019	TGCTGAGGCGCTTGAAGTTATTTCCCTATACTCTGGCTGAGAATGCTGGTTTGAACCCGATT GCCATTGTTACCGAGCTGAGGAATCGTCATGCACAGGGTGAGATAAATGCTGGAATAAATG TGAGGAAGGGTCAGATTACCAACATCTGGAGGAGAATGTGGTGACGCCCTTCTTG[T/G] AGCACAAGTGCGATCATGTTGGCGACAGAGTGTGTGCGGATGATTTTGAAGATTGATGATA TTGTAACGTGTGAGGTAGAGTTGATTGAACAATGTGGAATTTAGAGAGAACTGAATCACCCA TTATCTTCTTTTTTTTTTTTTTGTACTTCGTGCTTTAATTTTATGTTGGTAGTTT
>Gm05_3 8596037	GTAGATAAAATAGAGGTAGAGGTGTGGTAGATTTGAGAGCAACATTGAAATGCATGCCAAG ACAGACTCGGAGGTAACAAGCCTGGATGCGTCGTCCAGCACAAGGTCTCCTCGGCGAGCA GTGTACTACGTTTCAGAGCCCTTCCCACGATGGGAGAAAACGACGACGTCTGTTGCACT[C/ A]AGCCCTGTTCTCAGCCCCATGGTTCCCTCCTCACTCTCACTCTCCTCCAGCCGCTTC TCCGCTTCACGCCACCGCAATAACCATAATAATAAATCTTGGAAGGGCATCGACGTACG AAGAAGAGGGTCTTCTCCAATCCGAATTAGATCGCCAACATTCTCTCTCTCGTCGATA
>Gm05_3 8913395	AAACACAGAATATAAAAGAAACAGAAGGATTATATATCATATGATAGTTGAAACTTCCTTCCA AGGGAAGTCTCTCTCTCTCTCTCCCTCTCTCTCTCTCTCTCCCTCTCTCTCCCTCCCTCT CTCTCTCTCTCTCTCTCTATATATATATATATATGTACATGTCTGTGTGTCTATA[A/G]TAAGT TTCTAACATCTAAGGATAGCCCTGTCTGTCTGCCAATCCGAGGGGTGAGCGAGAGTGAA GGTGGTTGACATCTCTGAGAAGGGCACGCCTCATGTGTAGCTGAAGTGAAGGGTGCTTTAT TTTCTCAATTTCTTCTTAATTTTGATCACTTGGCTGTGAACTGCAAAACC
>Gm05_3 8913666	CACGCCTCATGTGTAGCTGAAGTGAAGGGTGCTTTATTTTCTCAATTTCTTCTTAATTTTGA TCATTGGCTGTGAAGTCAAAAACCCATGCAACCCTCTCGCGGAGGAGTAGGGGTCTGT TCTCCATCTTCTTCTGCAAACTCACAAACAGGACCTTTACAGGAACCAATATGTG[C/T]GA AATGTAATTTTGGCGTTAGGGATACCTATTCAAGGATATATATATATATATACCCGGAAGAA GGAAAGAAATATCAGATTTTGAGAGAGAGGAAATAAGGAAAAAAAATCTTTATTTATTATTA TTATAAAAAAAGGAAGTTTATATGGGCTCCTTCCCGTCCACCTTCTCCATT
>Gm05_3 9082457	TACATAGAATTACTTAATATTTGATTTTTTCATGAGAGTGATTATTATTGCTGCAGCATGTG GTGCTAGTGGGGCTCTGTATACGCTGATTTGCTGTGTCAATTGGCTGTGGCTGCCTATACTC TTGCTTCTACCGCCCCAAGATGAGACGACAGTATGGTCTAAAGGGAAATGGTTGTT[C/T]GG ATTGCTTGANTCATTGCTTCTGCGAGCCCTGCGCCCTCTGTCAAGAATATCGTGAGCTTCA ACACCGTGGAATTTGACATGATTATTGGTATGCACTCATTATATATATATTAACACTACTCTAT TTGCTAAATACCCAGCTATGCAAAGTTGGAAAACAAATTTCTTATTTTATAGA
>Gm05_3 9082469	CTTAATATTTGATTTTTTCATGAGAGTGATTATTATTGCTGCAGCATGTGGTGCTAGTGGG GCTCTGTATACGCTGATTTGCTGTGTCATTGGCTGTGGCTGCCTATACTCTTGCTTCTACCG CCCCAAGATGAGACGACAGTATGGTCTAAAGGGAATGGTTGTTNGGATTGCTTGA[C/T]TC ATTGCTTCTGCGAGCCCTGCGCCCTCTGTCAAGAATATCGTGAGCTTCAACACCGTGGATT

	TGACATGATTATTGGTATGCACTCATTATATATATATTAACACTACTCTATTTGCTAAATACC CAGCTATGCAAAGTTGGAAAACAAATTTCTTATTTTATAGAGAAACTCCTTTT
>Gm06_5 721969	TTTTTGGTAGAATAATATCAACATTGACCAGAAGCATAGACACGATATTTCCCGCCATCAAG TTAAACAGGTACGCTCATTTATTTTAACACAGCCCTATTTAAATATCCACAATTTTGTGCTTC ATTTCCCTTTTATTATTTCACTGTTGGGGCAGTGCTAATCAGTAACTCTGAAATA[A/T]NCAG GTGATATGTTCTCTTTGTGAGACAGAACAAGAGGTGAGTAGTGAGTTTAGATTTTCTTTAGT ATTTTTGTATGTAATCCTCCACAATGATATGTTAATTGTAAGGGAAGACTTAAATACTTGATT AAGCTGGATTAAATCATTATTTTTTGTCAAATAATAGAGAAATGATCTCT
>Gm06_5 721970	TTTTTGGTAGAATAATATCAACATTGACCAGAAGCATAGACACGATATTTCCCGCCATCAAGT TAAACAGGTACGCTCATTTATTTTAACACAGCCCTATTTAAATATCCACAATTTTGTGCTTCA TTTCCCTTTTATTATTTCACTGTTGGGGCAGTGCTAATCAGTAACTCTGAAATAN[A/C]CAGG TGATATGTTCTCTTTGTGAGACAGAACAAGAGGTGAGTAGTGAGTTTAGATTTTCTTTAGTA TTTTTGTATGTAATCCTCCACAATGATATGTTAATTGTAAGGGAAGACTTAAATACTTGATTA AGCTGGATTAAATCATTATTTTTTGTCAAATAATAGAGAAATGATCTCTG
>Gm08_3 461787	CAAAAACAACGACTGCACATTGTGCCCTCACGAGATCACCATCTGAGGTGGATCCTCAGCA GCTGAAAGATCTGTCAATTACAACCTTAGATACATATTCAACACTCCACACCCATTTTGTATT GCTCAAAGGAATAATTTATGTATATCAAGGAAGAGGAGTTTAAATTAATTCAATGT[A/G]TATT TTTTAAGGAGGGACGCTGGTTATGATTGTAACAGGAGGTAAACATACATATGCTAATCACAT TTTTGAGCCAAACAAATTTATAGAAATATCTTAATTACAATTGGATGTAGTTACAATTTTCTAC TGCTATTGTTTTTCCGAAGTAGTGAACAGAGGATTGTGTTGCTGCATAT
>Gm09_6 112681	ATGCATTGAAGGAACAACAAAATAACTAGAAAGGTATTGAGTTCAGAATAACCTGTGAAATA AACGTGCATATGGTTCTAGAGCCGGTAATCCAGCAACGAGATGCTCATCCAAAGCTGTTGT TGGAGGAGGAAGTAGGAGTGCCGGGACAGCAGCTGCTGTTAAAGTAGCAGTCTCATA[A/T] CGAGCAACTTCATCATCACAAACACTTAATATAACACCAGCACCATTAGCAACTGCCATCT TGGAGTTGATGGCAGAGTTGAGGATCTTCCCAGAACCAGCAGCAGAACTAAAGAAAA ACAAATTCAGTTAAGAAAAGATTATTATAAATCTTGCTTTTCGAGAATCAAAATGAT
>Gm10_3 8131569	CGTGCCATGTGCGCGCTCAACTTTATTATGTGTAATGTGGCTTTTGGCAGTGGGTTGTTCAA GGAGCAGTCAATGTAGATATTGGTGCTAACCTTCTGCAGAAGGTGGGGATGAGGACGAG GGAGTTGATGATCAAGCTGTCAAAGTAGTTGATATTGTTGACACTTTTAGGCTTCAGG[T/A]A NATGATGGAGGGTTTCTTATTTCTTGTTAACCATGAGTTGATTTCTTCTAAAGAACAAATAAC TTTATTATTTATTTTTATTTTTCAGGAACAACCACCTTTGATAAGAAACAATTTATTACTTAT ATGAAGAGATATATTAAGTTGCTGACGGCCAAATTAGAGGGTGAGCAGCAA
>Gm10_3 8131571	TGCCATGTGCGCGCTCAACTTTATTATGTGTAATGTGGCTTTTGGCAGTGGGTTGTTCAAGG AGCAGTCAATGTAGATATTGGTGCTAACCTTCTGCAGAAGGTGGGGATGAGGACGAGGG AGTTGATGATCAAGCTGTCAAAGTAGTTGATATTGTTGACACTTTTAGGCTTCAGGNA[A/C]A TGATGGAGGGTTTCTTATTTCTTGTTAACCATGAGTTGATTTCTTCTAAAGAACAAATAACTT TATTATTTATTTTTATTTTTTCAGGAACAACCACCTTTGATAAGAAACAATTTATTACTTATAT GAAGAGATATATTAAGTTGCTGACGGCCAAATTAGAGGGTGAGCAGCAAGA
>Gm10_4 7783606	CCCTTTCTTCTTGTTCTTCCCTATGAATATTGTTTTCTAGAGTTCACAATAGCGTCCATCCT GCTATCTGGTATAATTTGTAGGGAGGTGGCTCCATGCCACTATAATATAGACTATAGAATA ATTAACTTTTTTTTTTATGTATTCATAGTTTTATTCAACTAGTCTTCTGCGG[T/G]TGCAGC TAGTGACCTGGGCATTGCAAGATAAACAATGGAGGTTGTTGGCATGAAGCTAGGAATGGA CATGCATATTCTGCTTGTTGCGTTAGTAATTACTAAGTTCTTGAATGTTGACACTTTACTCTG CATGGTTTGTAATCTATAATAGAACTAAATTGTGACATAAATTTTCC
>Gm12_3 862466	GGTGGTGTAATCAAGAAGGAATCAATTATTACAACAACCTCATCAATGAGCTATTGGCTAA TGGTCACAACCTCACTATTTCAAACCTCTGTTTTTAAATTACATTTAAATTTTAGCATTTAATTA TTAAGTGTTGTCAACAAAAATTAATCAAAATATTGTTTGTGATTGAGACAATTAC[T/A]GGTGTA CTGCCATATGTGACTCTTTTTTCAATTGGGATCTTCCCCAAGCTTTGGAAGATGAGTATGGTGG TTTCTTAAGCTCCCATATAGTGTGAGTCTCCATTTTATACTATCCAGCAAATTTAATGTTTT TCTCATCATCAGTAGATGTTAGATTGTTAGTTTTTACTAAAAATGTTT
>Gm12_3 8459068	TTGGATGGAGTGAAGAAAGAGAGAGAAGCAATTCGGTCCAAAATCAAGCAAATTGATGATG CACTGAAAGCCATAGACAAGGATATCCAGTCTCTACAGGAGGAACTGACAGCTGTTTCTCA GAAGAGGGACAAAGCTTTTGAAGATATACAGCAGCTAAGAAAAACAGCGCAGGAGGGG[A/ G]TATGTTTCACTTTAAAGTATGAGCAGTGCAAGTGATATTTAAGGGTGTGCTTGGTCTGA ATAGGTGGAAAAATGAAAAATGAGGAATACAGGGAATGCAAAATGGAGTGAGATGGAATGCAA CATATGTTTTTGATTTTCATCTGGTGTTAGGCTGAGAAGGTGGAGAATGGGGAGGTTT
>Gm13_3	AGTGATCAAATGAATAGTGAAACTCTGATTCTGATTGGAGAGTGATACCAAAAGTAATAAA



5524268	GAAACTTCTGGCTCTAAGTATTGTGCTTTTTTTATTTGTGAGGCATTTTGGTTATTGCATTAT GTGTTATGCTGTAATGCATGCAGGCTGCAAGGAATGCTGGCGTGCCAGTAGTGTT[T/A]GAT GCGGGGGGCATGGATGGGCCCTTCCACCACAATTATTGAATTTTGTGATATTCTGAGTC CTAATGAAACTGAACTTGCTCGCCTTACCGGAATGCCAACAGAAAGTTTTGAAGAGATTGC ACAGGCTGCTTTGAAATGCCATGAATTGGTCAGTTCAAGTAATGACTTTAGGGG
>Gm13_3 5617464	GTGATAATTGCTGCACGGAGCAACAACGAGATGGATGCATCCAACTGAAGAACGAGTTCC CTGAGCTTCTTTCCATCAAGGAATCACTGATCAAGTATGTCTTTGAGCCAAACAAGAAAAC GCTTAAATATATTAAGCGGTTTCATGTTTTAGACATCACAGATTATGCATTTTGT[G/T]GCG GGCTACATAGATCACTCATTATTTTCATTGTTATTTTGTGGGTTTTGATAAGAACCATACTAC AAAAGCTAGCCATGTAGTTCTTGTCTCTTAGCTATAAGCGATTCTGTTCAGCATATTTCC ATTATATTACCGTGTTTAGAGGGTGACAAGTATTATAACTGGGCCAGATA
>Gm13_3 5823484	AGCCTCGTCCCCGACGTCATCGATCTCAGACCTCAAAAAAACTCAAAAGGGTCCTCCTCAA ACCTCCCAAAGAACACAACCTTTGGACCAGATTTCGATCCAACAAAGTTAAATGCCTTCTGG GTCGGTCACAAATTGAAAAGGGTGTGTGTGTTAGAGAAGAAGGTTGTGTATAAG[A/G] AGAATGTGGAGGAGATGGTTGGAAGTGNCAAAGTCAGCACATTTGGTGNATTTGGGAAGT GGGTTTGTAGCCGGAGAGTGATCGGAGATCGGAGGAGATCTGGGTGCGAGAAATGGGAA ATGGTTTTGATGAGAGTGTGTGTGGATGTGTGTGAGAAAGAAGCGGAGGAGAAGAAGAAG
>Gm13_3 5823512	GACCTCAAAAAAACTCAAAAGGGTCCTCCTCAAACCTCCCAAAGAACACAACCTTTGGACCA GATTTCGATCCAACAAAGTTAAATAGCCTTCTGGGTGCGTCACAAATTGAAAAGGGTGTGTG TGTGTTAGAAGAAGAAGGTTGTGTTATAAGNAGAATGTGGAGGAGATGGTTGGAAGTG[G/A] JCAAAGTCAGCACATTTGGTGNATTTGGGAAGTGGGTTTGTAGCCGGAGAGTGATCGGAGA TCGGAGGAGATCTGGGTGCGAGAAATGGGAAATGGTTTTGATGAGAGTGTGTGTGGATGT GTGTGAGAAAGAAGCGGAGGAGAAGAAGAAGAAAAGGAAAAGAGTGAGGAGGAGGAGGC
>Gm13_3 5823533	GGTCCTCCTCAAACCTCCCAAAGAACACAACCTTTGGACCAGATTTCGATCCAACAAAGTTAAA TAGCCTTCTGGGTGCGGTACAAATTGAAAAGGGTGTGTGTGTGTTAGAAGAAGAAGGTTGT GTTATAAGNAGAATGTGGAGGAGATGGTTGGAAGTGNCAAAGTCAGCACATTTGGTG[G/A] ATTTGGGAAGTGGGTTTGTAGCCGGAGAGTGATCGGAGATCGGAGGAGATCTGGGTGCGGA GAAATGGGAAATGGTTTTGATGAGAGTGTGTGTGGATGTGTGTGAGAAAGAAGCGGAGGA GAAGAAGAAGAAAAGGAAAAGAGTGAGGAGGAGGAGGCACTTGATTAGCCAAATTTGCG
>Gm13_3 5823811	GTGTGTGAGAAAGAAGCGGAGGAGAAGAAGAAGAAAAGGAAAAGAGTGAGGAGGAGGAG GCATTGATTAGCCAAATTTGCGGATCAAATTCATGATCACGGCGTTTGGATTTGTATAGAG AGAGAAAGAGGGAGTGGGTGTGGAGATTTAGAATCCAAAAAACATGCCGCCGCCGCC[A /G]CCACCACCACAGCCTAACCTTTCCCCCCTCACTTTCTCTCCCCCTCTTCTTGCTTAT GTCAGTGAGTGTTGATTTGTGAGAGAGCCTTCACAGTTCACACTTGTGTTGTGGAGTATCA CTCTGTGACGACTTTGATTTGGTTTTCTTTTTGGTTTTCTATTCTTTTTCTGAATT
>Gm13_3 5835159	TAGTTGCCGTACATTGCGAGGCGGAGAAGATCTTCGCCGCGATGGCTCCTCCTCCTCT CTCGCCTCTTGTTGTTCTCCCGCTCCTTCCACGTTGGCTGCCTCGCGACGGACGTCATCCT CTTCTCGCCGAAAAATCAAGAAGCTCCGAAAAGAACAAGGAACAAAAAAGAAAC[A/G] CACACACGCGAACACGAATGTTAGAAAGAAACACAGCACACAGCGCGAGTAAGCGAATTAT AGAGGAAACGACATCCTTTCTGCTTTTTTATATAAGGTGAGCTAAAACGACGACGTCGGAT TTGAGAGATTTGGGATTGTGATCTGAGTAGATCTAGGCGAGGGAGGTTGGGTGGTG
>Gm13_3 5862124	ACTAATTAATGTCTTATGCTACCAAAGCACTCATACAATCACTAACAATGACGAACACTCAA AATCACTAGTCTTAATTAACATCTATCAAATATGTTTTCTGAACTTTGCAATTTCTAATTC TCTAATTGCCCAATTTCTTTGCCGCGTTAGCTTCAACCTAGCCTCGTCATTTCT[T/A]TTTGC TTCCGTTGTTACCATTCGGCATTTTTGAACCACCTCATTATGTTGTCCACCTTTCCCTGG CTGATGGGTGACCNCTTTTCCCTCCAACGCTTCCATGGCTGGGGCCTGACCCTCTTGTTT CTCCGTGCGTTCTCACTCTAAACAACAATAACGTCAGTTCTTAATTCATAT
>Gm13_3 5862205	AACATCTATCAAATATGTTTTCTGAACTTTGCAATTTCTAATTCTCTAATTGCCAATTTCTT TGCCGCGTTAGCTTCAACCTAGCCTCGTCATTTCTNTTTGCTTCCGTTGTTACCATTCGGCA TTTTTTGAACCACCTCATTATGTTGTCCACCTTTCCCTGGCTGATGGGTGACC[C/T]TCTTT TCCCTCCAACGCTTCCATGGCTGGGGCCTGACCCTCTTGTTCCTCCGTGCGTTCTCACTCT AAACAACAATAACGTCAGTTCTTAATTCATATTTTATATATATATGCAGAACATCTAG CTAAGGCTGAATGAGATTTATGCATGTAAGAGTGTTTAACTTGATCACT
>Gm13_3 9517326	GTAAGAGACAATGTGTGTTCTACAAAAGGTCATCTGCCTTTCGTTTGAACCTTTGGGTGTTT CTTCTGTTGTAGGACAAAGACAATATCCCCTGTGACTGTATCAGGCTGCAAAAACAAAAATA GCTCAAAAATAGTTTTCCATAATCTGGTACAGTGTCAGCATCAAAGCTCAACATA[C/G]GCG TTCATCAGCTTCACCAGGGAATGTAATCTTCTGTCCATTCTGCATTCCCTTCTCCACAACAA

	CTTCAAGTACTTTCTTCTCCTGAACAACCTTCTCTCCCTTGCACTGTTGGCACCTGTCTCTG TCATTGATTGTCTCTCCAGTACCCTTACACTCATTGCAAGGATGCTGCATTT
>Gm14_2 8088505	ATTGTAGTTGTTACTTCACTCTAAAATGCCAGCAAGCATTTTTGCTTCTCCTTCATCAAGCAG TGATTTTCATTTTATTTCTATTAAGATAGCTATAGCATTCTGATATCTGTTTGCTACTTTGCTCT GAAGTCTAACATAATTTTTTTCAGCTAAACATTTTTTCTCTTTTCTTCTCTA[A/C]AGAATACA AGATCATCTCAAGAATTCCCAAACCAATCTGCACAAGGAGGATCAAGAAGCCATGGCGAGC CACCTCCATTTTCTCTATGCCGTTTAACCCCGTGATATTGCAAGCATGTTTCATGAACCTC ACGAACGCTACAAATGCACAGCAGCAGGGGTACATTCTCAGGAAAGG
>Gm15_3 100509	CATGAATAGTAATGGTTACTATTTTTCTGAGACACTCAAAGGAGATCTTGACAAATTCAGGCC ACCGTTATATTCCATTTTAACCCGTGCTGGGAGCTGAGTAGCCCTAGAAGTTGCGGCAATA TCTCCAACCAAACCTTCAACACTTTCACTTTGAAAAGGGTAGGAATTAGAAACAGCT[A/G]AA GTTGACTCAAACCAATTCAAGAAAGGAACTGCTGGAACCGCCACTGCTCCACTCCTCCTC CTGCAGTAGCAACACTGAAGTACCATTGCCATGACCTCCAATTTGAAACCCCATTTGAGAA CCGTTCTGTGCTTGAATCTCACGCAAGTTAACACCTGTATTCCCTACACCATAG
>Gm16_3 0464934	TCTCTCCCTCCTCATGAGGAGAAAAAGAAGCTTTCAACTTTCTATTATTCATCCAAACACC ATCTCAAATCGAAAGGAACAAGCTTCACCATCATCATCACCTTTTCACTCCTCATGAGTA CCAACCAAAACACCACCTTCTTCCCCCTCCCATGGAACAAATACTACTTCTGAA[T/C]TTG TTAATAGCCAGAACAATCCATCCAGGCTACAAATCAAGGTAACTATATATTATTGCCAGG TTGATTATATATCATATTATTCTAATAATTCATATTACCATCGATGTTATTATATGGTCCTCTC AATTATATATATATCATATCATATCATATCAATAATCTGGATCCACATCT
>Gm16_3 0479527	GCGTGGAAGCCAGTGTTTCATCAGGCGCCGAACCGAGGATGATATCAAGCTGGCAAACAG TGTTGATGTTGGATCTCTTCGCGATCCCCAGCTGGATGCTGATAGAGTCAAGGACCCAGAA TGGCTCATTGTGATTGGGGTTTGCACCCATTTGGGTTGCATTCCATTGCCAAATGCTGG[T/A] JGACTTTGGGGGATGGTTTTGCCCTTGCCATGGATCACATTATGATATTTCAAGGCAGAATTA GGAAGGGGCCAGCACCATACAATCTGGAGTTCCCTACTTACACCTTCTTGGAAGAAAACAA GTTGATGATTGGTTGAAAAATAATATGGTTCTGCATACAAGCGGAATGAGTTTACAA
>Gm16_3 0539483	GGTTTGCGGAACTGCCAGACTGTGTTGTGCCAACCAACAGGCGGACGGGCAAGGTTAAC CGAAGGGTGCTCTTTTAGGAAGAAGGGAGATTGAATGGTTTGTTAACAGACTTTTCTTTGT CAATGATTGAAAATTTTCGGTTGTTAGGTTTTGAAATGGAATCCATTTCCGTTATGG[T/C]AT ATTAATTTTGTTAAATTGTAAGACAATATCTCNNTCAGCCTTCCACAAGACATAGTAGCCAAT GTTGAATTTTTCTTTATTAAGTGGATGTGGTTTTGTTGACAACATTTACTAAAATCAATAT GGTTTAGTTTGATGGTTATTTTACAATAACATGTTTTGTATGCTCTGATG
>Gm16_3 0539518	CAACAGGCGGACGGGCAAGGTTAACCGAAGGGTGCTCTTTTAGGAAGAAGGGAGATTGAA TGGTTTGTTAACAGACTTTTCTTTGTCAATGATTGAAAATTTTCGGTTGTTAGGTTTTTGAA ATGGAATCCATTTCCGTTATGGNATATTAATTTTGTTAAATTGTAAGACAATATCTC[T/A]NTC AGCCTTCCACAAGACATAGTAGCCAATGTGAATTTTTCTTTATTAAGTGGATGTGGTTTTG TTCGACAACATTTACTAAAATCAATATGGTTTAGTTTGATGGTTATTTTACAATAACATGTTT TGTATGCTCTGATGACGTGTTAGTGTCATAATGTACATATGAATTTTG
>Gm16_3 0539519	AACAGGCGGACGGGCAAGGTTAACCGAAGGGTGCTCTTTTAGGAAGAAGGGAGATTGAAT GGTTTGTTAACAGACTTTTCTTTGTCAATGATTGAAAATTTTCGGTTGTTAGGTTTTTGAAA TGGAATCCATTTCCGTTATGGNATATTAATTTTGTTAAATTGTAAGACAATATCTCN[C/G]TCA GCCTTCCACAAGACATAGTAGCCAATGTTGAATTTTTCTTTATTAAGTGGATGTGGTTTTGT TCGACAACATTTACTAAAATCAATATGGTTTAGTTTGATGGTTATTTTACAATAACATGTTT GTATGCTCTGATGACGTGTTAGTGTCATAATGTACATATGAATTTTG
>Gm16_3 0626524	TATAATGAGGATGGTGTCAGGGAAGAGGTACTATGGTGAAGACTGTGATGTGAGTGATGTA CAGGAAGCAAGGCAATTTAGAGAGATCATTAAAGAGTTGGTGACGTTAGGAGGGGCTAATA ACCCTGGGACTTCTTGGCTTTGCTTAGGTGTTTTGATTTTGATGGTTTGGAGAAGAG[A/G] CTAAAGAGGATTAGTAAGAGAACCAGATGCGTTTTTACAAGGACTCATTGATCAGCATCGTAA TGGAAGCATCGTGCCAATACTATGATAGATCATCTTCTTGCTCAACAGCAATCACAACCTG AATATTACCCGATCAAATCATCAAAGGACTTGCACTGGTAATTAATTAATTATT
>Gm16_3 1191863	CAACCCCAATCCAGTGATGCAGCTCATCTCCCCTGCATCCTCCACTGGTGAGAACACGCAG CACAACGCTGCCAACACCTCCAAGGCATCATCGGATTCCGAGAATTTGCTGAGTCGGTGA TCAAGGCTCCTAAGCAGGCCTCTGGGGAGCACAAGAAGAAAAAGAAGATCAAAGTGAC[G/ T]TTCCCATCAGGTCAAGAGCGGAATGCACCATCACAGGCAATTAGGAAATGCTTGCACTGT GAGATAACCAAGACACCACAGTGGAGGGCAGGGCCAATGGGGCCGAAAAACTCTGCAAT GCTTGTGGCGTGCGCTACAAGTCAGGCCGGCTTTTCCCCGAATATCGCCCTGCAGCGAG
>Gm16_3	ATGTACTACCAAGCAAGGGGTACATGCCACCTAAGTTGGAATTTTCATATTTCCAGGAATC

1204160	AAATTGTTAGTGATTGATATTTTTAGTGGTCAAAATGAATATTTAAAAATAAAAAATAAAAAATAAA AGTGAAGGAAAGCATGAGATAGATAATAGAGGAAACAACGTGAGAGTTCCAC[A/C]GGA GTAAAGTAAAGCTTTTGTAGATTGAGAAACCTCACTTGTGTGGTGTGGTTGTGTTTTT GCATGCTTCGGACATGTCGATGGCGATGGTCCACCCTTCGGTTCACCCAATCGAAGCCCC ACCCATGACTGAGCATGCAATTGCAATTGCAATGCCAGACACACTCTGAAGGA
>Gm16_3 1204451	CCCAATCGAAGCCCCACCCATGACTGAGCATGCAATTGCAATTGCAATGCCAGACACACT CTGAAGGATACGCAGGGCATGCCCGGCACCCTCGGTGGCTTCCTCTTGCCTTCGCCAG TTCTCTTCGCCCTCGTTTCCCTCTCCGTCATGGCTACCACCTCCGATTTCCCTTCTGT[C/T] ACTGCCTTCGGTATTCATTAATTCCTTCTTCTTCTTCTTCTGTTAATCGTGCTATTTTT AATTGGGTTTGTGGTAATTGTTGATTTGGGCTTGTGTTTTGTGTTCAAAATCAGCACCCAG ATTTATTTTTTTTCAAGTGGGTTTATCAAGGGAGCAATGAAATGAACCTT
>Gm16_3 1225684	TAACAAAACAGATCATATATCTGCCATTATTCTAAAACAGAAAAGGATAGTTAAAAATAAGAA TACAAGGTTACTCCAGAATAACGATGTAGTAAAAGGAAAGTATAATGTACAAAGGTTTCTAA ACAATACAATATGGTGAAACAAATAGGATACCTGCATAATTTTCAGGTGCATCAGG[C/T]CCA CACATTGAGGCTCCGAGAACCCTTATCTGTCTCAGCATCAACAACAAGCTTCATAACAGTTTT TTCTTGTGCGCTATTTGAAGAATATCAGCAACCATAATCTAGATAAGAAAGTTATGCTGCAA AGAACATAACTAATACTGGGTCCAACCTGAGATTGAAGAGCATGTGATGATAA
>Gm16_3 1461554	TGCAGAAAATGAAGGAAGATGTCATGCTGTTGCAACCTGCTTAACCAGGAAATTATCTCTTC CACCCTCTACATTTCACTCTTTTCTTGACCATCATCATCTCCAAAGTGACCCCAACCT ACTGGTTCAAGTAACAACCTTGATACCTGGTACACCGCGCTTGTACGAAGCCGTGC[C/T]GT GACAAGGGACAGAGTAAGAACTGGAATTTTGATGAAATTGCAATGGAGTCCTAGTCTAAA CTATAAGATAGAATAAGACTAGGGAATTTTCTTGTGGGATAGTTTATAGCATGTTATTGCTAT TTTGCCCTTTGAATTTGTTATGGGTGCAATTTATCTCTTCTCTAGTTTTT
>Gm16_3 1475163	TATAAGTGTTGTTTTCAAATATTAAGGAGTTGTGGGGTGCAGCGTGATTTTCTCGTTATAA AAATAAGAAGGAAGAGTGATTCAAATCTGAAGCTGAACCCTAACTAGCTGAGCAGACGGAG GAGAACTCGACTGAAGCTGCCTCTGCGACGCCGTTTCGATTACGCCTCAGACATTTT[C/T]G TAAGTTCATCGTTTCACTTTACACAAATACCCTATTCTTCGCATACTTCGTCGCTGCAACC CCGAACAACGATAACCTAATCAATTTGGCTTCGCCCCCTCTTTTTCTCTTCACAAACAATAA CAATTCACGTGTGATTGATTTCTCCAGTGTTTACGAGTTTCAAAGATGCTCG
>Gm16_3 1476359	CATTTATCAAAGCCCAGGTTAGCAATGCAAAACAGTGAAAGTGCTGCTTGCTGCTGTAG GTTGCAGTCATGGTGCAAGAAAATGGCAAGAGGAAGAAACCAACATTCTGGTGACTGGCA CACCAGGGACAGGAAAGACAACCTGTGTGCACTGCTCTAGCTGAAGCCACCCAGCTCTG[T/ C]CACATCAATGTCGGAGAATTAGTCAAAGAAAAGAACTTGCATGATGGATGGGATGATGAG CTTGATTGTTACCTTCTTAATGAAGACTTGGTAAGCTATCAATTTATATAATTTGGATCATT CTATATGCTTGTGTTTGAGATTTAGCTCCTCTAAAGTAAATGAGGAGATGTGTAA
>Gm16_3 1827645	GCATGTGTTGCAACTTGTTTACTGCAACTCTTTCTGTATAAATACCATATACACACAATGAA ATGAAAAGCTTGAGTTTCTTTCTTTCAATTATTCTAACAGAGGGTCGATCTAAAGTATTTATG GGAAGATCATGATTAGGGCATTATTTCTTAAATAAGCAAGAGAAAATGGACAA[A/G]ATAGA AGCACATAATACTGAACACTTTATCCTTATTACATGAGTCATATGTCAAACAGGAAACAACA ATCCCCAAAATAGAACACACCCATATCCTTAGCAATTAACAACACCACACAAGAAGAACTA CACATAACTTACAAACTTTTGCTCCATTCAAACTGATCAACCAATTGCA
>Gm16_3 1827884	GAAACAACAATCCCCAAAATAGAACACACCCATATCCTTAGCAATTAACAACACCACACAA GAAGAACTACACATAACTTACAAACTTTTGCTCCATTCAAACTGATCAACCAATTGCACCA CCTCTTCCACCACAGACGTCAACAAAGACTTCCTAGCATTACCACATTGCACTTG[C/G]CA CGAATTTACCTTGATTTCCAGTGAGCACATCCAATTGACTCAACTTTATAGTAGCATCCAC AAACTTCTCAAAGAACAACGTTTGGTTCAACGCAAATGCATTNACCAACCCCTTAGTCCTCT TATCATTGAGCAAGTCCTGGTCAGAGGTGAACACCCCTTGCGGTTTCATTAGG
>Gm16_3 1827991	ATCAACCAATTGCACCACCTCTTCCACCACAGACGTCAACAAAGACTTCTAGCATTACCA CATTGCACTTGNCACGAATTTACCTTGATTTCCAGTGAGCACATCCAATTGACTCAACTTT ATAGTAGCATCCACAACTTCTCAAAGAACAACGTTTGGTTCAACGCAAATGCATT[G/C]ACC AACCCTTAGTCCTCTTATCATTGAGCAAGTCCTGGTCAGAGGTGAACACCCCTTGCGGGT TCATTAGGTCAAGGTAGTATTTGTTGTGCAACACAGTCGGGGTTCTGATGTCCAAGTTGAC GGTGTGCGCGAGTTTGCGTNGGGGCAAGTGGATTGTAGTTGCTTGGCTAGGGT
>Gm16_3 1828137	CAAAGAACAACGTTTGGTTCAACGCAAATGCATTNACCAACCCCTTAGTCCTCTTATCATTG AGCAAGTCCTGGTCAGAGGTGAACACCCCTTGCGGTTTCATTAGGTCAAGGTAGTATTTGT TGTCGAACACAGTCGGGGTTCTGATGTCCAAGTTGACGGTGTGCGCGAGTTTGCGT[C/T] GGGGCAAGTGGATTGTAGTTGCTTGGCTAGGGTTTTGTCCATGTTGGGGTCGAGAGGGGA

	GAGCCTGTTGAAGAATGTGCCGCAATGGGCACGACCAAAGGTGTGTGCGCCGGATAAGGC GACAACGTCGGTGACATCAAAGTTTTTGGCGGCGAAAGCGTCAAGGGTCACGCCGGTGG
>Gm16_3 1829738	ATATCACAAAGAAAAGCTTCAAAACTATAGTTAGTAAGAGCAGAGGGTTTTGAGAACCAAAA TAGTAGTGAAGTAAAGGTAAGAGGGATGAAGAGATTACAAGGAAAACCTGAATCACGTGCT GCCAAAACAGTGATATCTGCGCAAGAAACGATCCTTCCACACTCCTTATGGATAATA[G/C]C TCTGATGTCATCAATGGTTTGCAAGGCCTCAGTTCTGATGCCTCCGTTGGCTGGTTGGTCT CTTTCACCTGGACTCCCATCTAGCAGCAATGATCCATCACATCCCTATACAAATGCAAATTA AGTTAGAGGTAATATCTTTTTTCTATATATATATATGCATGCATATATATTAACC
>Gm16_3 1840753	GTTGGGGGACTCGGCATTGGGACACGTAGCTATCAAATTGTTGTTGAAGTTTGGGTGCGATT GGTGGGTCCGTTTCGATGGTTCTGTTCACTAGGGAGGGACAGTGGGCACGACCGTAGGTG TGTGCGCCGGAGAGAGCGACCACATCGGTGGCATCGAAACCTCTATTTCCGAATCCTCT[C/ G]AGAAGATCGTCGGTTCGGAAGAATGGTGCCGGTAGGTTGTCCGGCGCGGTGGCGTTTG GTCCTAGNCCGTCTTTTCTTCCCAGTGGCACGTCAAAATCAGGGCCTCCTAACTGTTCAAAT TACAAAGAAGTTACCAAAAGGAACAAACGAGAGATCAAGCATTATGTTTAAATGTGTTT
>Gm16_3 1840819	GTCCGTTTCGATGGTTCTGTTCACTAGGGAGGGACAGTGGGCACGACCGTAGGTGTGTGC GCCGGAGAGAGCGACCACATCGGTGGCATCGAAACCTCTATTTCCGAATCCTCTNAGAAG ATCGTCGGTTCGGAAGAATGGTGCCGGTAGGTTGTCCGGCGCGGTGGCGTTTGGTCCTAG [G/A]CCGTCTTTTCTTCCCAGTGGCACGTCAAAATCAGGGCCTCCTAACTGTTCAAATTACA AAGAAGTTACCAAAAGGAACAAACGAGAGATCAAGCATTATGTTTAAATGTGTTTTTAATCT CATTATAATGTTATAATTTGAGTGAAAATTATCTCTAATTTATATATTATAAATGATTT
>Gm16_3 1842815	GAGCTACTCCACTGTCCCTTCTCGAAGACATCCTCAAGATGCTTCCTTATAATCCTCTCAAGC TTGGGACATGTCAACAAATAGTAATTCATGATAGTCCAGGGACTAGTTTCCCAGAGGAAA CATGGATTTGAGAAGCCAATACAAGGGAAGAAATCAAGAGCAAAGAGTGAAACAAAG[G/C] ACTAACCCTAGCCATGTTTGATTAATTTGCTATGAGTGTGACTTGAGATAACCAAAACCTTAG AAGAGTTATTTATAGATTCAATATTGAGGTAACGTTCTTCGGTTGCTTCCCACTTCCCACCC ACTAGTCTTCTACGTATAATATTTAAAACGTGGCGCAAGCGTTTTTTGAGATTTG

**Supplemental Table S2.** SNPs identified by Method 2 on the single-library comparison

This table contains 255 SNPs identified using Method 2 on the single-library RNA-Seq comparison between the Clark/IsoClark 10-day root experimental data. The file contains a unique SNP identifier that includes the genomic location. The sequence surrounding each SNP based on the soybean Glyma1.01 genome assembly is also provided.

SNP	[Clark/IsoClark]
>Gm01_2 205875	TGTTATGCCTAACATTCAACCTTCTTCTTCCCAAGAAGGCTGGGGGCTCATCCAAGGG TGCTGCTGCTGACGATGACTCTTAAATAGGGATTCACTGTTACAAAGTTCTTGAATTTAG ATTCTATGTTTGCTCTCTTTTTTTTTTCCTTTTATTTGTAGTTATGATAATTGTGCC[A/C]TT TGTTAGTAGTAGTTAGGTTTTTCAGTCAGATGTGTTTAAAGAATGCTCTTACGCATCAAGAAA GGTTGTTCAAAATTTAAATTTATTTAATGTCACCTTTGAGTCTTGATCTAGCATTTGATTTCT AATTGCATTCTATCAAACCTTTAATTTAAACTGTTTGATTAGGCTTTGAGAGATT
>Gm01_6 704379	TTTGTTTTGTGTTTGATTTTTTTTTTCCTTTTCTTTCTTTGCTGTGAAATTTACCTCATTTTCATG ACCCACCCCTTTTACTTTTTTGGGGTCTTAGCATTAACTCTGTGTTGTGCTGTGCTGTGC TGTTGTGTTAGACTGTTAGCACCTTTTTCCATAGGCAAAATCATGTGTTACTCTC[T/C]CAAG CCTCACTGCAATAACAGTTGTTTGGTTCTGCTTTTGTTAATTGCCAACAACAATC ACTTAAAGCTTGCTCAGGAAAAACGCATACAACGCATTCTCTCTCTTCACTCTCTTTCT CTCTCGTACTTCATCGTATATTTAATGAATAAATTTGTAGTAGTAATAAATT
>Gm01_5 0660171	AGCAAGTTTTGAAGTGCACGATGTGGGAGTGAACCTGCTACTGCTAGTTGATGCATGTT TCCTATGAGAGGTAGCTTTTTTGGACCAGGGGGAAGTTTGTGAGCAACATTAGGTTTTAG CTTGAGATATTTGCAAGCCAATTCAGTAGAATGAAGAGGGATAAGGCAATAACCAAGA[G/ A]GTAGGTTTGAGCTTCCATATATGGCTTTATTATAGAGTCAGCTCTCAAGGGTTTTATTGT TGTTTGTCGATGGCTGGTTCCGTTCCATTTATAGAATCTTCAATCCGCAAGGAAATTTG ATCTATCTATCATTTTTTAAACCACTCAACGTAAGTCTTATATTCTTTTCAAATATCACTT
>Gm01_5 0660172	GCAAGTTTTGAAGTGCACGATGTGGGAGTGAACCTGCTACTGCTAGTTGATGCATGTTT CCTATGAGAGGTAGCTTTTTTGGACCAGGGGGAAGTTTGTGAGCAACATTAGGTTTTAGC TTGAGATATTTGCAAGCCAATTCAGTAGAATGAAGAGGGATAAGGCAATAACCAAGAA[C/ T]TAGGTTTTGAGCTTCCATATATGGCTTTATTATAGAGTCAGCTCTCAAGGGTTTTATTGTT GTTTGTCGATGGCTGGTTCCGTTCCATTTATAGAATCTTCAATCCGCAAGGAAATTTGCA TCTATCTATCATTTTTTAAACCACTCAACGTAAGTCTTATATTCTTTTCAAATATCACTTA
>Gm01_5 3617124	TATTGATACAAAGTTATATTTTTTCAATTCATTCAAACCTAGCGAACTATGCATCGGTTCCA AAGTTCACATTGGCCCCGACGCAGGCACGCAGCTCAGTTAGCAACAGCGTTTTTTTTTTTT TATCACTTCATTCCCAACCCTAGCGAACTATCAATCGCTCCCAAAGTTCACATTGG[A/T]GC CCGACGCAGTCTCAGTCCGTACCAGCGTTAGTGCCAGCTCCACTGTGCGGTAGGGCTCG GAAAGCTCCGCATTTGGCCATTCGCGCTCTGTAAGTTTCCCTTTCTCTAATTCTCTCTCTC TCTTTTCATCAAGCTTTGTTTTGCTTTGGACGTTTTCTCTTTGACCTAAGATCCAAAGT
>Gm02_9 54265	TGATCAATGAAGAACTCCAGAAGCCATCAAGGTTGTCCATGCCTGTGGATGCTTTCCCT GGTTTCTTCACCACAATAATGGGAGTCAAGTCTTTTGTCTCTTTTTCTTTCCACATTGGACT CTGTGGCACTAGATCATGGTGACAGCATATTATCAAGGCTTCTAGTAATAGCAAGAT[A/C] GAAGCCATTATTAGTAGACTAGAGATGGGACATTTGATTGGAAGGCTTTGCTAGGAATAG ATGAGAAATATCATATATATATATAAAGTTCTGCCCTTACCCTTACATGTTTTTAATATCAA AATTTGAAGGGAAACCTTCTTTCTTTTGGAGAAATAAAAAATCTGGATTGCTTATGA
>Gm02_6 692680	TCAATCATGTTAGAAAACTGAAGTCATATCTTCAAACGATGCTTCTGAATTTCATACACG TTTACCAATTTTAATTCGAAACTTGTGCACAGGTAAGAAACCAAGCAATAGTGAAGTGGAT CAGCTAGTGAATCTTGCAGAAAAATACACTCTTGCAAAACCATCTATTTTGGGGCTT[T/C]TG GGGACTTATTTCCGTATGTTTTTTTTTTTTGTTTTTGGATAAAACATGTTTTTTATCTCTC AGAATATCTTGAAATTCAGTTTTAGTCTCTATAAACATTAAACCACTAGGGTGGGGTTTGG GTGGTATACATGAAGGCTTAGGTTTCGATCCCTAATACAGTATGTTTTTAGTTTC
>Gm02_1 7023049	TTTCCATTAGATTTAGAGGGACTTGAGAATTTAGTCTTCTGTTTTCTTTTGGGTAATTGTG TGATTGGTTAATGCTTGCAGTGAGGAGGCTGCAAGGAAGCTTTGTTGTATAGTTACAAGT CTGCAGCCAGTGGGTTCTCAGCTAAGCTTACTCCGGAGCAGGTTGAACAGATTTCAG[A]

	GTAAGTCTGATTGTTTTCAATCTAATTTTTATTATTCATTTCCCCCAGCTAAACTCCCTG ATGTAATGTAATATGGTTCATTTTAATTTTTTTTTATTTTATCCTTTTGAAATTCCTGATGC ATTGTGGCTCTTTTATGCTGATTGTTTATGTTTGCATCTAAGCAATATTTGTGTT
>Gm02_1 7023050	TTCCATTAGATTTAGAGGGACTTGAGAATTTAGTCTTCCTGTTTTCTTTGGGTAATTGTGT GATTGGTTAATGCTTGCAGTGAGGAGGCTGCAAAGGAAGCTTTGTTGTATAGTTACAAGT CTGCAGCCAGTGGGTTCTCAGCTAAGCTTACTCCGAGCAGGTTGAACAGATTTCAA[C/T] TAAGTCTGATTGTTTTCAATCTAATTTTTATTATTCATTTCCCCCAGCTAAACTCCCTGAT GTAATGTAATATGGTTCATTTTAATTTTTTTTTATTTTATCCTTTTGAAATTCCTGATGCATT GTGGCTCTTTTATGCTGATTGTTTATGTTTGCATCTAAGCAATATTTGTGTTT
>Gm02_4 2350182	CGTGGGTGAAAGCGACCACATCGAAGAGCTTTGCCAAGAGCTTGGCGTCTGTTGAGGTGCG CCATCGACGATGAAGACGTCATGCTTGGCGAGGAGGGACTTGCGGGCTTTTTTCAGAGA CGGATCGTAGTAGTCGTTGAAGTTGTCGAGTCCGACGACGCCGTCTCCGCGTCTGTTTTAA C[A/T]CGAGGGAGACGTGGGAGCCGACGAAGCCGGCAGCGCCGGTGACGAGGACGGAC ATGCCGCTTGACGGTGGATCTGGGCGGAGGTGCGGACCTGCTTCTCCCACTGGATCCC GCCCCAGGAGGCGGAGAGGTAGCGGCTGCCGAATCAACGAAGCCTTGGAAGCTGAGA TATGAAGCGG
>Gm02_4 4574156	CTTCTCAATTTCTACACCCTTGTTGTTGTTCTGGTTTTTCATAGGCTGAAATGTGCACTCCAA GTGCTGCTCTACCAGAAGGATCAAGGTTGTTAGACCAAGCAGCATCCCACTCAACAGCTT TGGCTGTGCTACATGCAACACTTGCTCCTCCAGGAACGGTGAGCCTAGCTGAGTTCT[A/T] CAAAACCAATCAGCAAACACTTATTTGAATCAGCCCTTATTGAATTACATATTCACATTG ACTAGTGATATGGTCAAAATAGTGTATATAAGTCGGGGACAACCTCTTATCTTATGAGCTAC CTTTCAGGATTTTGTTTGGCCTAAATCCAAATTGTAAGAAATTAATTAGGAGTTGG
>Gm02_4 4574157	TTCTCAATTTCTACACCCTTGTTGTTGTTCTGGTTTTTCATAGGCTGAAATGTGCACTCCAAG TGCTGCTCTACCAGAAGGATCAAGGTTGTTAGACCAAGCAGCATCCCACTCAACAGCTT GGCTGTGCTACATGCAACACTTGCTCCTCCAGGAACGGTGAGCCTAGCTGAGTTCTG[A/G] ]AAAACCAATCAGCAAACACTTATTTGAATCAGCCCTTATTGAATTACATATTCACATTG ACTAGTGATATGGTCAAAATAGTGTATATAAGTCGGGGACAACCTCTTATCTTATGAGCTAC CTTTCAGGATTTTGTTTGGCCTAAATCCAAATTGTAAGAAATTAATTAGGAGTTGGT
>Gm02_4 8476266	CAGAATCATAGAATCAAACACGACATGATCTTCAAAGCAAAATTTGTCCATTTGATACCA AAGCTTTAAACAATGGCATTGGATAGTTCTAGTAGCAAGAAGATAATATCTTTAAGAATGC AGAAGACTTTATAAAGCATCAGTAACCGAGGGAGGTTTGTGACTTATGACAAATGG[G/A]G ACCGAAGAGCTCTGTGTGCTTCCTCCTTCTGCTTGCCTGCTTGTACATTTCTCTGTCTC GATAACAATAACCTCTTGACCTATATTAATTACAAAGCTTACCAATGTCAGTATACAAGGA GACAAGTAATTTCAAGCAGCAAAATGGACTTGGGTAGAGCTTTTGCTAAAACCTGT
>Gm03_3 5154151	CATTTGGAGAGCAGCTGTGGTTCATAAACCGAGAAACATTTCTTCATTTTTTGCTGATAT ATAAAGGGGGGATGGGATTTTCGGAGCTTCAGTATCACCAGGAAAAACTCCAGTTGCTG ATAAATGCGAGTAGAATCAAAAATGTAATCATCTTCATTGTCAACCAACAGCTCCTCC[A/C] CTCTAGCACTATCTATGACTTCCCCTGCATACTCACATATGAAAGTTTCTGCACGAATAGA ATCCCATGATCTAAGACCCCAACCTTTATTCTTGTTCTGAAAACCTCCAAACGAAATTTT AAGCCACTTTGAGAAACCCGATTTTCGGCAATTAGAAGGACATTGACAAGAAGGGCCAC
>Gm03_3 6460374	GACATTGCAAATTAATTTTTCCATACAAGTTTCCTCAATCCTAATATAGTGATAGTGATAA ATAGAATAAGACAACAAATTAGTACTAGGCTCTGTCAATCTTCATAGGCCAATACTTTTAC CACTCTTTAATAACTTATGACCCGATATTTGAAGGTGAGGATAACAATTGTAACA[T/C]GAA AATTACAGGAAATACGGATTTTATTACATGCACGTAAGAAACACATCAAATTTCCACTCTCA AGAATCCAAAGAAAAGATAACAATGCATGCTCTCAAATCCTCAACATGCTAACTAACATCT TGCCAATGCAACTTCCTAAGGATGAATATTTAGACAAAAAAAACATTAAAGAG
>Gm03_3 6554101	ACCCATGTCTAGATGCACTATGGGCATCAGTGACATCAATCGCTCTTTGATGTGAACCTCTC GTCCACTCTACAACCACCAATTATCATAGGTGAAGTCATTTTGTACATGAAAGTTAAAAAAT AAGATGTTAGGCCGATTTTATTTAATGTAATTGTGTGTAATAAAATACATATAAAT[G/A]CAA ATACATTTAATACTTTTTGTTATTCCTAAAAATTTATTGAAAGATCTAATATTAGACCTA ACATATTCAATTAAGATGTCATCAAACAATGATTTAATTAATTTAATATAGATAAAATTGAG AATTTTTTAAATAGAAGGATCAAATATATTTATTTTAAATAGAGAGACC
>Gm03_3 6559131	CATTCCAGCTACAGTTTCGCGAGTTTTAAAATACATACCTGAGTTACAACATCAAGTGCAA GCACTAACTAAGAAAAAGAGGAGCTTCTGTGCAGAATTTCAAAAAATCTCAAAGGAGATT CGGTGAACAAAGAATCTCAAAGGAGAATTTCCCATCACAATTCTGATTTTGCTGTTT[C/T]A ACTAGTAGGCTCAACGATTGTGAAGCTGTTGTTTACATTTCTCTTATGAGGCTCACAAGG CTCCACTATCCGACATCTTGCAATGTTTAGAAAATAATGGCCTTTATTTGCTAAATGCTTCT

	TCCTCTGAAACTTTTGGAGGAAGGGTCTTCTACAACTTGCATTTCCAGGTATCTTT
>Gm03_3 6559857	TGTTTCTGCAGGTGGAAGCAAGGATTTCTAATCAAACGATCTTAAGTGCTCTGATGTTAAAT GTCAATATATGAGAAGCAAAGGATTTCTAATCAAACGATCTTAAGTGCTCTGATGTTAAAT TTGACACCTAAAAGAGTCTTCACATATTTGTCATGCATGAATGGCAAGTTCGGAAA[A/T]TA GAATGGAGTAATCACATCACATGCCGCTTCTATATACATATGATTTGGTAAAAAAGCAAGT GGCCGGCAGGAAGGTGCATGGATTTTAGGAGGAGTTAAAGAAAAGAATATTGTATAAACA TAAATGTTACAATCAGCCTCACGTAGCAAAGTGTGACACCCTCTTTTCTCACAAATA
>Gm03_3 6559926	ATGAGAAGCAAAGGATTTTCTAATCAAACGATCTTAAGTGCTCTGATGTTAAATTTGACAC CTAAAAGAGTCTTCACATATTTGTCATGCATGAATGGCAAGTTCGGAAATTAGAATGGAGT AATCACATCACATGCCGCTTCTATATACATATGATTTGGTAAAAAAGCAAGTGGCCG[A/T]C AGGAAGGTGCATGGATTTTAGGAGGAGTTAAAGAAAAGAATATTGTATAAACATAAATGTT ACAATCAGCCTCACGTAGCAAAGTGTGACACCCTCTTTTCTCACAAATATATGTAATAA ATGAAAGGAATAGAAAGTTAGAATTAATGAAAAGTTTAAACACATTTAAATAAAA
>Gm03_3 6560002	CATATTTGTCATGCATGAATGGCAAGTTCGGAAATTAGAATGGAGTAATCACATCACATGC CGCTTCTATATACATATGATTTGGTAAAAAAGCAAGTGGCCGGCAGGAAGGTGCATGGAT TTTAGGAGGAGTTAAAGAAAAGAATATTGTATAAACATAAATGTTACAATCAGCCTCA[A/G] GTAGCAAAGTGTGACACCCTCTTTTCTCACAAATATATGTAATAAATGAAAGGAATAGA AAGTTAGAATTAATGAAAAGTTTAAACACATTTAAATAAAAGCATTTCAAAAAGGATAAA AGGTTTACATTCTGTTTTCTAGCATTCTAATAAAACTTATTTAAATAAATAAATAAAA
>Gm03_3 6560077	TATGATTTGGTAAAAAAGCAAGTGGCCGGCAGGAAGGTGCATGGATTTTAGGAGGAGTTA AAGAAAAGAATATTGTATAAACATAAATGTTACAATCAGCCTCACGTAGCAAAGTGTGACA CCCTCTTTTCTCACAAATATATGTAATAAATGAAAGGAATAGAAAGTTAGAATTAA[G/A] GAAAAGTTTTAAACACATTTAAATAAAAGCATTTCAAAAAGGATAAAAGGTTTACATTCTG TTTTCTAGCATTCTAATAAAACTTATTTAAATAAATAAATAAATCATCTTCACTCAAAATCAG GTCGTCCAACTGAATGAAAAGAAGTCAAACCTAATAGTAGAAAACATAAAATAAC
>Gm03_3 6952394	AAAAGAAATAAAAAAATAAGATCTGTTTTTTTTTTTTTTGAGAAAGATCATTTTTTATTAT TATTGTTTTGAATTTTGAGAATTATGTAATGATGAGAGTTGATGTTGTGAAAAAATAAT GGAAATCCAGGGAGGAGGCGGTGGTGGAGGTGGCAAGAGTGGAAAGACAATGT[T/C]CA CCAACAGCAAGGAGAATTTGTTTCATAGTTTTTCTTGTACACTTTGGTACTCTTCAAACATT GGTGTGATCCTTCTGAACAAGTATCTGCTCTCAAACATGGAATTCAGTTCCCAATCTTCC TCACAATGTGCCACATGTCGGCCTGTGCTGCTCAGCTATGTCTCCATTGTGT
>Gm03_3 6959955	AACGAATTGCAAAAGTTTTGGAAGTCTTCTCAAATAAAATGCTCCTCAAACAGCTGGTTC TGACATAAACAGATATAAACTAGTTATGTTCAAGTGCAGCAGAGAGGACTGTTGTTGGCATAT TTGCCCTCAATCATTTCTTTACTCAACACCCATATATGTTTTGTCATGCCTGGAAAT[C/T]TA GTATGGTATCAAGTCTTCTCAAACACTACATAGAATTAGAATGCGGTAACGAATGGCAGCAT TAAAGCACACACTACACTAGCAGAAGAAATCAGCAAAGCAAGCCGGTAATGCCCAAAT GTAAAAAAGTAGCTTGGTTAGGCAATTCATGTCAATGTTTCATCTACTTGTTTTTCG
>Gm03_3 6996777	CATATATTTGCATTCACCTAACGTAGTCTTGATACCCCAATCCACTAACAATATTTTTTAT ATTATTATTATTATCATCATCAAAATGCCTGGCTTGTAATGAAATGGATTTGGACCAA ACTATCTAGTGACGTGCGCAATGAAATCATCCAGTTCGCGCGAGAAACACCC[C/A/G]TTC ATCCTTGGACCTGCGAATAGCATTTTTAGGTTTCATTGCCCTTTGTCTCATCTCATCACCTT CTTTTGTGCCATCAATCTCCTCACGGCATTCTCAACATCTGATGATGTCACCAACTCGTC CCTGTGGTCCCAGTCTTCAACACCCCAACTCTGAGCACTTCTGTTACCAA
>Gm03_3 6997184	GCACCCCATTTGTCATGCTCTCCATGCAAGAGTTCCATCCACAGTGACTCATAAACCCAC CAGTTGAACTGTGACTTAGAATCTCCAATTGGGGTGCCAGTCTCTCACAACCAACCCCG TGCCCTTCACTCTCTCTTCAAACCCCTTTGGAAGCTCAGCAGTTCTCACGCCATCCTCA[G/ A]GGAACACATCTCCCTTGTGAGCATCCCTCACCACCCATATGAACTTTTGCTTGCTTTTT CCAACCCGTTTGCAACCTCCTTGATTTGCTCCTCAGAGAAGCATGTTGTTGTCCCAAACGA CACATACAAAAGTACCCTGCTTCTTGTGTTGTCAAGCCACTCAACACTGAAGTGCCTTG
>Gm03_3 6997185	CACCCCATTTGTCATGCTCTCCATGCAAGAGTTCCATCCACAGTGACTCATAAACCCAC AGTTGAACTGTGACTTAGAATCTCCAATTGGGGTGCCAGTCTCTCACAACCAACCCCGT GCCTTCACTCTCTCTTCAAACCCCTTTGGAAGCTCAGCAGTTCTCACGCCATCCTCA[T/ C]GAACACATCTCCCTTGTGAGCATCCCTCACCACCCATATGAACTTTTGCTTGCTTTTTTC CAACCCGTTTGCAACCTCCTTGATTTGCTCCTCAGAGAAGCATGTTGTTGTCCCAAACGAC ACATACAAAAGTACCCTGCTTCTTGTGTTGTCAAGCCACTCAACACTGAAGTGCCTTGT
>Gm03_3 7024958	GAAATGAAGAATGGGGTTTAGAGTTAGTCAAGACAAGTGGTGTGGTGTCTATGTTATGA CTGCACACATGTGAAGTGAAGTAGAGACTATTAGTCCACAGCAGCTGTTTCTAGTGTGT

	GTGTCATTGCATTCTCATCCTTTTCTCTTTTTTACGCGCTTAATTTCTTCTCTTT[A/T]C TCCCTCTTCTCTCTGGAATTTGGAGCATCAGCCAGCACTCTATGGATTCTCTGATTGGTA ATTGGCCATCCTACGATCCTCACAACCTTCAGTCAGCTTCGACCTTCCGATCCTTCTAGTTC TTCTGTAAGTTGCTGTTGTTGTTGTCTATGAAATTGATAATCCTGGTAATAATTACT
>Gm03_3 7027198	GTAAAGCAACCTTTTTGTAGAAGGGTTCTTCTATAATAACTCTAGTTCATTTTCTCCAATGT TATTTTTCAAGTTGCTTAACCTTTGTTTTTTAATTTTCGGTAGACTCAATGGTGTCTTAAT TCAAAGTTTGCTAACCTGTTGAAGTTCCAACAGTGATAAGTACTGAAGCCAAAA[C/T]CATC CTCATGAGACATATTTATCAGCATTCTGAGCAGAAGGTTAGTGAAGTTGATGGTTAAAGAGC ATGTGTTTTGGTGCAGTTAGGTATGCATATGCTTGATGCTCATAACCTTTTGTGTTTTTCAG TTGAATCCAAAAAGAGCTGCATCTGATAACCTTCTTTTACCAGAGCATGGATG
>Gm03_3 7144398	TCCTTAAAAAGTTGACACCAAACTGAAAAAGCATTTATCTATGTTGCAAATGTTTAGGGCT TTCGTAATAATTGCAACAAGCAGATGGGAGTTTTTCAATGACAGGTTCAAGAAAGGTGAAA GAGAACTTCTACATGAGATTCTGGCGAAGAAAAGCTTGGACCAGCAAGCAACAACCA[A/G/T] GGCACCACCAAGCAACACTCCAAGACTCTGATGAAGATCAAAGGTCCTCATCAATCTC ATCATCTTCTGGTTACACTACCTTGTGGATGAAAAACAAGCGACTCAAGAAGGAGAATGG GGTGTGAACTCTGAGCTCACAAGTATGAAAAGGAAGTGCAAGGAACCTCTTGATTTGGT
>Gm03_3 7165409	AAGATTCACACGTACCATCTCGAAACAGTTGTTCTGATAACTGTTCTGAGAAGCTCCATC ACAGACAGGTGTACCAGAGATTAGTGTCTTACTCTTCTTCTTATTGTCAAGCCAGCGCCCA ATTAAGCCTATCATTTCTCGTGTCAAGCTAATCTCCGCAATGGATGATTTTCTGAT[A/C]T GACGGAGTCAGTGCCCCCAATGCAAATTTCAAGGTCATTAAGTAAGCTTCTAGGATGAGA AGGGACATCAAGAAGTCTTTCTTGAATAGCTTGGCAGAGGTTTTGCAAGTCTGCAGAACT CACTTCAGTAGGTTTGACCTGGAAACAGGGAAACAGAGAATTACATATTTGAAGATGCT
>Gm03_3 7827399	GTCGAGTGATCCATACCCGCTAGGTTTTACACATTCTTTTCCCTGAAACAAACACCACCCC TTTTAGGTCTTCTTAATCCTCCCTCTCTCTAACTTTCACACGCAGCGGCTTCTTCTCCG TTTTCTGAATCGGTGTGAACAATGGCGACGTACGCTGCGATGAAACCCACCAAGC[A/C]A GGGTTGGAGAGTCCCAGGAACAGATCCATAAGATAAGGATCACCTTTCTTCCAAGCAC GTCAAAAACCTCGAGAAGGGTATGCACCTATTTATTCTACTTTTTGTTTTTAAGGTTGTTGC TATCCTTAGCCTTATGTTTCTAGTTTTGTGTTACTGTTAGTTTGTGCGGACTTGGTT
>Gm03_3 7828684	CAACACATGGGATAGATTTGAACCTCGTGTCCACAAGAGAGTGATTGACCTCTACAGTTCC CCAGATGTGGTTAAGCAGATTACCTCTATCACAATCGAACCTGGTGTGGAAGTTGAGGTG ACCATTGCAGATGCTTGATCTATATCCAAATTTTAACTGTTATGGTTTCTAGTTTTT[T/A]G TCTCCTCTTATTCTGTATTTTATAAGTTTTGATTGCGGGACCGAGTTGAATTGTTTTCTTTT GTTTTACGGTTGTAGACGGTTGATGAATACCTGAGAAATGTCATTTTTATATATTCTAGAAT CCTACATATATTTGTTTTGAAATTATTACGAATCATTTCTAATGGACTCAGCTAA
>Gm03_3 7828791	TGGAAGTTGAGGTGACCATTGCAGATGCTTGATCTATATCCAAATTTTAACTGTTATGGTT TCCTAGTTTTTCGTCTCCTCTTATTCTGTATTTTATAAGTTTTGATTGCGGGACCGAGTTGA ATTGTTTTCTTTTGTGTTTTACGGTTGTAGACGGTTGATGAATACCTGAGAAATGTCA[C/T]TTT TATATATTCTAGAATCCTACATATATTTGTTTTGAAATTATTACGAATCATTTCTAATGGACT CAGCTAAGATTATATCTTTTCTTTTCTTTATCATGTTCTGCAAATTTTAAACCTGAAACT GATTGTGAAGTAGATAAGTTGGAAGGTGTTTTTGCAAATTTGTACAATTT
>Gm03_3 7832228	CCTTGTGTTTCATGAAACTGCCTTTTGTGTTTTAGGTACCAACACATGGGACAGATTTGAACT TCGTGTGCACAAGAGGGTGATTGACCTCTACAGTTCCCAGATGTGGTTAAGCAGATTAC CTCTATCACGATTGAACCTGGTGTGGAGGTTGAGGTGACCATTGCAGATGCTTGATCT[C/ T]GATCCAAATTTTAACTGTTATGCTTCTAGTTTTTTTTCCCTCTCTTTTTGTATTTGTAAG TTTTGATTGTGGGACTGAGTTGGATTGTTTTCTCTAGTTTTACGGTTGTAGACATTGATGAA TACCCGAGAAATGTCATCTTAATATATTCTAGAGTGTTACATATTTTGTGTTGAAAT
>Gm03_3 7863675	CTTTGATCTAATGTTTGCAAGAGCATCAGTCCCGCTAACGGTTGCAGCATCCGAGTGCCA CGCTTCGGGAAACGCCCTCATCAGCTTGTTGATACCCATCCAAACAGCCAGCATCAGCTCCCC CTTCGCCTTGTACCCCTTTCTGTCTCCAGCCTATACCACTGCGGTGCCAAGGGACTGT[C/ T]TGGTGGAAACACGCTTTGGGATCTCATTGAGGTCAAATAAGACACGACCAATGAAGTCAT CTTTCACGACATCCTTGTCTTTCACAGTGACCTCCAGTATGGAAGCCTGAATGCGGTCTTT GGAGAAAGCAAAAACCTGGTTCATTGCGGATTAGACTTCTTGTGCAAGTGCCGAGTGGT
>Gm03_3 8065066	TGTGGAACCACTCAAGACTATGTTACCATAGAGATCCTTCTGATGTGCAGATCACAC TTCATGATAGAGTTATATGTTGTCTCATGGATACCAGGAGATTCCATCCCAATCATGGATG GCTGGAAGAGAACTTCAGGGCACCGGAATCGTTCAGCACCGATCGTAATCACCTGCC[C/T] JATCAGGTAGCTCATAGCTCTTCTCAACTGCTGAGCTGGTCTTGGCAGTTTCCAACCTCTG CTCATAATCAAGAGCAATGTAGGCCAGTTTCTCCTTCATGTCCCTCACAAATTTCCCGCTCC



	GCAGATGTGGTGAAAGAGTAACCACGCTCAGTCAAGATTTTCATCAGGGCATCAGTGAG
>Gm03_3 8083417	GTTTTCTTTTCTTTCCATCCATGGATTCTCTGAGTTCCATCAACGGCGATTTGGGCTTCG ACGGCGGCACGGATCGCAGATTTCGCGTTTTTCGCGCCAAGCCTCCTTCCAACAACCTCAC ACGCCGATAGACATTCCGGCGCACGGCCACTACGACCACCACCACTACTGGTCCCCGCG [T/A]GACGATAAAACCTCCGCGCGCTCTCTCCAAATATCCTCTCTCCTCCTTCGTTTTCT CCGTTTTCCGAAGCGTTAGGTCCGGCCACAGGTACATGAAGAGGCTCTTTCTCATGATCT CGCTCAATGTCGCGTACTCCACCGCCGAATTGCTCACTGGACTCTTCACCGGTGCGGTAG GT
>Gm03_3 8117453	GTACATGAAATTATAATAACCGTTTTCTGCACTTGTACTTAATTATTGGCCTTACTTGCTAT TCTCATGAGCTACAGCAGCAATGTGGTTGCTTGTGGCCCCAGCAAAGTCTTATGGAAC GCGCTTTCACCATGCCAAGCTTCATCCTCTTCTTCGTGCGACGTCATTATTATCCCTG[G/A]C TTCTTCTCCTCCTTCTCCCCAAAGGGCTCAGCATCATCAACACACTCAATTTGCTTCTCGA GAGAACAAAAAGAAGGGGAAGAAGAAGAAGGGTGAGGCACGTGGTGCACGTGCACCTC CTTTGGTGCAGTTGATGCAATAATGGAACGGCTTCTAACACTTGTATGCTTCTGAGTCT
>Gm03_3 8117485	TTGTACTTAATTATTGGCCTTACTTGCTATTCTCATGAGCTACAGCAGCAATGTGGTTGCTT GTTTGCCCCAGCAAAGTCTTATGGAACGCGCTTTCACCATGCCAAGCTTCATCCTCTTCT TCGTGACGTCATTATTATCCCTGGCTTCTTCTCCTCCTTCTCCCCAAAGGGCTCA[G/T]CA TCATCAACACACTCAATTTGCTTCTCGAGAGAACAAAAAGAAGGGGAAGAAGAAGAAGGG TGAGGCACGTGGTGCACGTGCACCTCCTTTGGTGCAGTTGATGCAATAATGGAACGGCTT CTAACACTTGTATGCTTCTGAGTCTTCCCTTCCCCAAGTGATGCTCACACAGCGAGT
>Gm03_3 8132996	TGACAATTTACCTATAGAAAAATGGTTGATTTAATTACATAATTACTATGATGCTTATTAAGT AGTGGTTTTGTTTTGAGTGGAAAAGGTGCACATAGTTTTGGTCTGCGCAGTGTAGATTTT TCAACCAGAGACTGTTCAACTTCAGCGGCACAGGTAGCCCTGATCCCACGCTGAA[C/A]A CAACCTATTTAGCGACACTTCAGCAAAACTGTCCACAAAAACGGAAGCGGCAACACTCTGA ACAACCTCGACCCTTCGTCTCCGGACACCTTCGACAACAACACTTCCAAAACCTTCTCAG CAATCAGGGCCTTCTCCAAACAGACCAAGAGCTCTTTTCCACCAACGGCGCCGCCACA
>Gm03_3 8136913	TGTAATTGGATCTGGGGTCTTTCATGTTTGCGACGAAATTAGGGTTTACTCCCAATTTCCGCC CCAAATTTTGGGGGTTTTCTGTTTCTGATTGTTCAATTTTCATCACCCGACCCCGTAGAAC CTGGGTTTTCTCGATTTGTGTTAACTCGAGCTTTTTGGTGGAAAAATACTCGCTTT[T/G]C TTTTCCCTTTGTTGTTCCCTTTCCCTGCAATTGAAATTTGTAATAAAGGAAGCTTCTTTCTA GTTTTGTGATTGAAGCTGCGATGGAGGAAGAGACCTCCGATGCAATGAACCTTGATTTGA ATCTGGGCCAGGCCAGAACCCGAAACTGGGCCATAAGCAATGAAGCTGTGAACT
>Gm03_3 8170431	TTAGATATCTTAGTATGGGAAAATAAAACGAAAATGTGAAATCTTTCAGCTGTGGAAAGA TGGAGAGATGAAAGAAGAAGTCATTGGAGGCCATAAGGCCTGGCTAGTTATTGAGGAAGT CAAAGAAATGATCCAAAAGTATTTATGATTCTTCTATTTTATTTTTCTTGCTCCCTCA[A/G]C CCCCACCCACCTCACAAAAAAAAAAAAAAAAAGAAGAAAGAAAGATTACAATTGAAGAAA CTAATCTTCCAATGTTTTTACTTTTTTTCTTTAACATTTTTTCATGTTGGCAACCGCATGTAG CTGTGTATCATATATATATATTAATGGAAGTGTGTACCATAGCTGAAAGAAATTTA
>Gm03_3 8173719	CGTCGTTGTCGTTGTCGCCCGGAGACGGCGACACGACGACGCGCGCAACCTTCTTG GTTCCAATTCGTCTTAACAGGACCTCCTTTGGCCATGCTCTGCACCCAAACCTTAACCTT CTCCAAGATCAAGGGCTTCCCTTGGAAACCATGGCATCCTTAATCCTCTTTCCAACCG[A/ T]TCCCTCGTCGTCGACAGTAACCCTAACCTCGGGATCCTCGTCGGCGTTCTCGTCGGAG ATGTAGGGGATCTCGACGGTGCCGTGACCTTGAGAAGAGAGGTTCCAGGGTGTCTTT GGCCTCTCCCTGCCAATTGAGAGTGAGGCTGATCTCGTAGCCGGGGATGATCTTCCCTT TCG
>Gm03_3 8173815	GCTCTGCACCCAAACCTTAACCTTCTCCAAGATCAAGGGCTTCCCTTGGAAACCATGGC ATCCTTAATCCTCTTTCCAACCGGTCCCTCGTCGTCGACAGTAACCCTAACCTCGGGATC CTCGTCGGCGTTCTCGTCGGAGATGTAGGGGATCTCGACGGTGCCGTGACCTTGAGAA[ C/G]AGAGGTTCCAGGGTGTCTTTGGCCTCTCCCTGCCAATTGAGAGTGAGGCTGATCTC GTAGCCGGGATGATCTTCCCTTTCCGACATTGATGTAGGCCTCGCCGTGAGGGAGC GAAGCGACGTCGTTTTGAGGAAGAGGTTACCCTCGCCGTGAGGATAGGGAGGTTAGTG AGGAG
>Gm03_3 8174150	CGAGGATAGGGAGGTTAGTGAGGAGGCTGGTGAAGAAGGTCTTGACCAGTCGAGGCAGT TGGTCTCGGACCAATGCCAGTTGTGGACATTGGTGCCGTGCGGGCGGTCTCCACGATC CACCGCTTGTGCCCCCTACCGTAACGAGCCATTGCAACTGATGCCTCTTTCTGGGAGAG T[A/C]CCTTAGTGATTAATAATAATTGGATTGGGGTGAGGGTTATGAACTATGATGATTATT ATTATATGTGTTGAGTGTGTGTGGAATGGTCTAGAATGAAAAGGAAGGGTCTAGAATGTTG

	ATGTTGCTTTGGGTACACTTTCTTCACCAACCAACCAAGCCCGAGAAGGTTCCCTAGGG TT
>Gm03_3 8174157	AGGGAGGTTAGTGAGGAGGCTGGTGAAGAAGGTCTTGACCAGTCGAGGCAGTTGGTCTC GGACCAATGCCAGTTGTGGACATTGGTGCCGTCGGGGCGGTCTCCACGATCCACCGCT TGTCGCCCTCACCGTAACGAGCCATTGCAACTGATGCCTCTTTCTGGGAGAGTTCCTTA G[T/C]GATTAATAATAATTGGATTGGGGTGAGGGTTATGAACTATGATGATTATTATTATAT GTGTTGAGTGTGTGTGGAATGGTCTAGAATGAAAAGGAAGGGTCTAGAATGTTGATGTTG CTTTGGGTACACTTTCTTCACCAACCAACCAAGCCCGAGAAGGTTCCCTAGGGTTTTCTT TT
>Gm03_3 8718231	GATATAAAGAGGCTTACCGTTGATTTTGATTGAGGTTAAGTTTATGAAATACTCATCAGAC GGTTGTGGATTGTCCGTGACGACGGTGTCCGCAACGGGGTTTACGATGAGTTGTGTGTA AGTAAGGTCAATTTTGAAGAGAAAAGAAATGAGGAGGCAGTGAGGCGAATATAGCTG[ G/A]CCCTGTGTTGGCTGATGAAGCTGGCAGGCAGAGAGTGAAGAACGAGGTGAGGTTA ATGAAGTGCTGATTTGGGCCGGGAGGGAGTAGTTGGACCGGCCAATGAGGCGAGGCC CAATGCATTGGCGGCGAGGCCCTGGAGAAGATGGGCCGTGGCGCAGGAGAAGATGAAG TCGGAGAT
>Gm03_3 8942920	TAATTTCTTTTTTTTTTCGCATTTCTATTTTTCTCCTCATTTCTTGGGAACCAACAGTTA GATTAATGACAGTTAGATTGATGAGGTACCTGAATCCTGACATAGTCGGTACGATTCCAAC AGAAGGCGTTCCCTAGCATCTGGTCAACGGTCTCTGAAACACCGTCGAGCGCAAT[C/T]TC AATCAGCATTGACGTAGAGCACTCGCCGGCATTGTTCAATTCTCTTCGCCGGAGCTCCGTT TCCGATGGAGAGAACAAGGAGATCCTCCACGCCGTTACCGACGGGAAGTCGCGCTTGT TGTGGAGGACGTGCGTGACCGCCGCCGCCGGCGGATTGTTCACTAAGCCGCCGTCG
>Gm03_3 8942927	TTTTTTTTTTCGCATTTCTATTTTTCTCCTCATTTCTTGGGAACCAACAGTTAGATTAAT GACAGTTAGATTGATGAGGTACCTGAATCCTGACATAGTCGGTACGATTCCAACAGAAGG CGTTCCCTAGCATCTGGTCAACGGTCTCTGAAACACCGTCGAGCGCAATGCAATC[T/C]G CATTGACGTAGAGCACTCGCCGGCATTGTTCAATTCTCTTCGCCGGAGCTCCGTTCCGAT GGAGAGAACAAGGAGATCCTCCACGCCGTTACCGACGGGAAGTCGCGCTTGTGTGGA GGACGTGCGTGACCGCCGCCGCCGGCGGATTGTTCACTAAGCCGCCGTCGACGGC GG
>Gm03_3 9788794	ACTGGGTGAAACGAGCCCGCAAAAAGAGGGGTATATGCGAGTGGAATATTTAGTTTCGAA ACTGTATTATGTTTCACTTATGCGCTGTTTTTTCTTTCTGTTCCGTTTCCTATAAATCCATG CCACACCCTTCACCTTATTTTCCGCACATCATATTACTCTTTCACATCATTTCTCA[A/T]TCT TCCTTGCTTGCACTCTCATCAGGTCAAGTTTCACAACCCCTTCTTCTCCCTCTCTCAATATTG TTTTGTTTTTTCTCTTTGGCATCATTATCTTTCTCATGTGTCACCCCATGCACATGCATCTC GCGACTCCATCACACCGATTCTCTATTTTCTTTTCATAACAAAATACTGAGATATAT
>Gm03_3 9788799	GTGAAACGAGCCCGCAAAAAGAGGGGTATATGCGAGTGGAATATTTAGTTTCGAAACTGT ATTATGTTTCACTTATGCGCTGTTTTTTCTTTCTGTTCCGTTTCCTATAAATCCATGCCACA CCCTTCACCTTATTTTCCGCACATCATATTACTCTTTCACATCATTTCTCAATCTT[T/C]CTT GCTTGCACTCTCATCAGGTCAAGTTTCACAACCCCTTCTTCTCCCTCTCTCAATATTGTTTTG TTTTTTCTCTTTGGCATCATTATCTTTCTCATGTGTCACCCCATGCACATGCATCTCGCGAC TCCATCACACCGATTCTCTATTTTCTTTTCATAACAAAATACTGAGATATAT
>Gm03_3 9790874	TTCAGTTGCATTCAAGGAAAGGTTTGATGCTCCACCATATGAGCTTGATCCAGCTGGCTCA ACTGTTGCACAAGCAGACTACATTTGGAGGATAATTGTTATGGTGGGAGCACTGCCAGCT GCGTTAACTTACTACTGGAGGATGAAGATGCCGGAACCTGCCCGTTACACCGCTCTAG[T/ G]CGCCAAGAACACGAAGCAGGCTGCAGCAGATATGTCTAAGGTTCTGCAGGTTGAGATT CAAGCTGAACCGCAGAAAGAGGAGCAGAAGGCTAACTCATATGGCTTATTTTCAAAGGAG TTCTCCGTGCGCATGGACTGCATCTACTTGGTACAGCAAGCACATGGTTCTTGCTTGATA T
>Gm03_3 9795164	TCTGCAGATGACAAAGCATGAAGAGGTAATTCATTAAAGGTAAAACCTAGTAGATTCTGAT ATGAAAATGAATTTAAGTGCTTAAACATTTCAATGTGCACGGTCTCTCAGCACTTCAAGAT CCATTTCTGTGGGGTCAGTAGCCTGGATTTTATAATGGACACCATCTAGCTCACGTC[C/A] AAACCTGTCTTTAGTAGTGGCATAGAGCATCTTGGCACGGATCCGAGATGTTGAAGGGGA CCTAATCAATATCATGCAATGAATAACTAAGAGTGATATCTCTGGCAAAAGCATAGAATGA GTAAGTTAAGAAGCATGTTGTTGAGGATACATGAGAAGGCTGGGATAAAAGAAGTTCAA
>Gm03_3 9795203	GTAACCTAGTAGATTCTGATATGAAAATGAATTTAAGTGCTTAAACATTTCAATGTGCAC GGTCTCTCAGCACTTCAAGATCCATTTCTGTGGGGTCAGTAGCCTGGATTTTATAATGGA CACCATCTAGCTCACGTCTAAACCTGTCTTTAGTAGTGGCATAGAGCATCTTGGCAC[T/A]

	GATCCGAGATGTTGAAGGGGACCTAATCAATATCATGCAATGAATAACTAAGAGTGATATC TCTGGCAAAAGCATAGAATGAGTAAGTTAAGAAGCATGTTGTTGAGGATACATGAGAAGG CTGGGATAAAAGAAGTTCAATCCAAGTTTGTTCTTCACAAGAACAAGAGAGGAAGACAA
>Gm03_3 9964048	CGGGCCATTCTGCACCGATCAATGGAAGAATTTCAACAACAAAGCGAATCGCATTTTACA ATTTTCGATGCGGATCGGCTTATGTTAGGGTTTGAGATCGAGAAGAAGAGAATTAGATCAT ACGCGGAAGCAGCAAGAAGCTGAGAAGAAATGCAATTGGCAAGCGAAGACTTCGACTTC[ T/A]AGTGAGGAGTGCTTGAGTGCAACTGAAGTCTGCTGCTGCTTGGTCTGCTGGC ATTAATTTTAGTTTCCAATAATATCCTTGAATCCACGTTTTAACTATGCCTATCTTCGCAAAA TCAGAAAATGTTTTTTTTTTTTTACAATAAAGTTTTAGACCAGGGTACTACTTGCATATT
>Gm03_3 9966061	ATTTTGAGAATATTACATTGACTGCAAACACAACCTCAAAATCCATTAGTGCAAATTAACATT CGTCGCTTCATTCTAATAACGTGAGCCAGATTTTTGATGTAAGTGGTTCACAACACAAAAGG CATAAATGGCTCAATCACTTGACACCAGGCACCAGCACTAGTTCACAAAATACTGT[A/C]TA CGCATGGAACAAAAGACACTTAAATTTAGAGCTTCTACATCACTCAATACTTTGGAGGTG GCTGCTGGGTTGGTAATAATCCAAAACGCTTCTTCAGAAGAAGTCTGCTGCTTAACTTTT ATCATCGGGTGAAAACCGGGCTGCAGTTTCAGGAAAACCGGTAAGTACAGAAAATG
>Gm03_3 9967974	TATCGTTTTATGTAAGTGAAGATACATGGTCACTGCAAAAAAATTGAAATACATGAAATCA TTGCCAGAAACATAGCTCGTCAGATGGAACAATAATGAGTCTAGAGTTTTCGGAAACATA CCTGATCTTTGATCGAAGGAAAGAACAGCAATAGAGGTAGAAACCTGCAATGGAGG[C/T]G AGAAAAAGGGTTTAAGGTTTTCGGATCCTCTGGTCTATACAAAGTTTTTGTTTCAACGAG AATGGGCCTGTCAATTTAGGTATTGGTTATGGGAGATTTTATTCTACACCGAGCATATTTT TTCCTGCACCCAGCATATTTATGGAAGAACCAAAAATACCTTACGTATAGTCTATAAG
>Gm03_3 9986148	TTGTGAAGATGCTGCATGCAATCATTTGTTGAAATTTTTATGGCTGGAGTTTTATTTATACT TTCTACTCGTGTGTTCTTAACTTTTTATTTTTATCTTGTCTCTTAAGTATCATGATCAAGAG AAGGGACGCAATTCGTCCCGATATGGCAGCAGTTCAAGAAGACCCATAATCTCA[G/C]CGT CAACGAGGCAAGTTCCCTCTGGTGATCATACTGACAGTCTGACTGGCCGGCTAACCTCAA GTGGAAGCCGACAATCTGCCACACATAGAAATATTCAACCTATGCACGAGACAAAACAAC CAACTTACACACGCTCTGGATCCACCAGAGGCAACCGTGATGATCCTCTGCGGAGTT
>Gm03_3 9993121	CACTATTTGAAATATTATAATTAGGTGAGAGGGGAAAACGATGATTGGGAAGGAGAAGGAG AACCTTTGGCGGTGGGTTTCTTGGTGAAGATGTTTCATGGCGATCGGTGCGTGTGTCACA GTCGAAGGCAATTTACGGAGAGACACGCAAACACAACCCAACGATGGTGGAGAGAAGAA[ C/T]GATCCTCGCAATTTTCATGTTTTATTTTTTTATTTTTTTTTATAATATATGTACTTTCT CATTCGACTATATATAATTTTTAAGATTTTCAAGTAAACAGAAACCGCATAGGTCTTGGTGGG GTATTTAAGCATATAGTTGACTTGTCTAATTACACTAAGATGCTTATATGTTTTTGC
>Gm03_4 0055679	CATTACCTTAAGACTATTTAGCCGAGAAGGTAATACTTGAAGGAATAAATTGATGGTTTA GCCTTATTTTTCTTTTTAATGGCTTAAGCTATGTTTATTTTTCTTCCAAATTTTATTTTTTG GTTTGACATCTAACCTATACTTTCTATTGGTGGCAGATGTAAGGCACATACCTT[A/G]CGTG GAAAACGTGGTGGCAATATATAGAAGCTTGGATGCCACTTTTTGTGCTAGTTTGTTCAGT CGCGTTCTTCTGGTGAAGCCTCTTTGTTTTTGGAGAGCTTTTCTGATTGATTGAAGAAAA AAAGCTATGTATTACTTGATGCTGATTTGATATTGTCTTTTAAATTCGTTGAT
>Gm03_4 0056070	AAACAATTCTTAATTCAACTTCTAGTAGTTTAAAACTGGCAAGCTTCCTCGGAACCTTAAC AATTTGAATGAGGTTGGGTTGGCCAACTTTCAAATCCAAGAGTGTGGTGCATGATAAA GCTGAGGAGCTTGAGTTGTCTTGTGGGAGAGGGAAGCCCAATCCAATGCAATCTTCC[T/C] AGATGGGACTTGGGGGCCGAGAACAGTGTGGAGTTATGGAAATGGATGTGAATCAAGTA ACACAATCAGCACCTGATAGCCCGCCCTTTTGTGATACCAAAGGTTTCAAGATAATGACTGC AGTGATCAAGATAGTGAGCATGTAAGGAAACATACTCTCTTATTTATGGTCTTCTGTTTGT
>Gm03_4 0131229	TTATATGTTAACTTTACCAAGAACAATAAACCCACATAAGCCAACCCAGTCAACCCCTGA ACAACATTTTGCAAAACAAGGTGCCCTCAACCTCATTCAAATCCATGCACTCACAAATCAC AATGAGTAAAGCAGACTATAAAAAAGAAATTAAGATCACAAGAAAAACCAAGAAAAC[C/G] CCTCTTATAACTATCAATTAATCTCATCAAAACACTTCTCACTGAGAAGTCAAGCAAAAACA AACTATATGTGCCAAATGCCAATATTGTCTGCAATTCACCTAAACAATATCTGAAACAAC TTCAAGAAGCAGAAACATTACATAAACTAAAAACCCAGCTTAAACAATAGCATTG
>Gm03_4 0132046	GATTCATATGACCCAGCATTATCTGCCAAAACGCAACAATAACAATTAGAAAACAGAA AAAAAAATTAACAAAGACAGAGATCAATAATAAGGAAACAGAAGAAAAAGTGAAATACCTT CGAAGTACATGTAAACACCGTCCTTTTCGGCGCCACGGCTTTCGCTGACGGACGATGA[G/ C]GGCGGGGAGCACCTTTTTACGGAGATCAGGCTTTCCCTTCTTGACGGTGGCCATAACC ATGTCACCCACGCAAGCTGACGGCAACCGGTTTACGCTTCCCTTGATCCCTTCACTGAT ATGATGTAGAGATTCTTCGCGCCGGTGTGTCGGCGCAGTTCACCGTCGCCGCCACCGG

	CAG
>Gm03_4 0154315	TTGGTGACGTGAATTCGTATGCTGGGGATCCGAATTCATCCTACAGCAAGCTCAATTCA GTGCAACTGGTGAGTCCAAGCCAGCTGGTGGAGCGGCTGATTCTAACGAGGCCTCTGCT GGGGTTGGAAGCACAGCTGCTGACTCTACTATGGTATCTGACTACAATTCATCTGTAAAT[ G/T]GTGGTGTTGCCGGTGCAAGTACAAATACTTCTGGACTTGAAAACGGGAATGCGTTGG AGAATGCTGATGGGTGCGGCTGATGAGAAACAGCAAGCAGATGGTTATGGTGTGCCTTTTT CTTTTACCTTTTTACTTTCAATGGAATTATCATGTTTTAAATTTTATTTTTGGACAATAAATT
>Gm03_4 0160427	ACTACTTGAGGTCTGTCTTCACTATTACTTTCTTGCTGCTAAATTCACCTCATATTATGT AGTTGGTGATTCTTCTCTTTGTATTCTAAGTGTCTTCATCTCTAATGAGATTCTTTAATTC AATGTTACATGCAGGCTATATTGCATTTTGAGGCTATTCAACCCCTTGCCAAAGC[C/A]CGTT GATATTGACTTTCTGGAGTCATGGGTTGTCAAATTCATAATGCCTAACTCAGAGAGTCCTG GTGTAGCAAGTGCAACAGAGCGTGAGGAACCTTTCTAGTATTTTTCTGGAGGTAATTCCTTG TCAGAATGGTGTTCTGGCATGTGTTCTAAAGGAGTTTCCTTAAGAAATGTGTTT
>Gm03_4 0172340	CAGACTACGGTTTGAGGAAGAAGCCAGGGAATATGTCAACTCAGTTCATATAGATGATG ATCCAGTGGACGAGTATAGTCTTCTGAGCACCAGCAGCAGCTACAAGAAGAACTTGAAA CCGAAATTGTGGAGGAGGAAACTCCCGTACAGGAGGCATCTCCACCAATTCATAGCAATTC [C]CACACACTGTCCAAGAACCCTGTTTGCTCTTGTTGGAAGAGTCCTTTGAGGAGCCTCC TAAGAAAACATATGCATCTATTGTATGTAATACTTTCTATACATGTTCTTTTTTTTTGGCAG ACTACCTTTTTTTGTGTGTAGTTATGTACTATCCATAAAAAAGGTACCACATGACTGGA
>Gm03_4 0179701	CCACCATTGAGCAGCACTGAACCCATCTTCACTCTCACTTTCCATTTCTTCATCTTCTCTTC TTCACCTCAAGCCATTGTGCTTGGCAGTAGAGAAAGAAGGAATGGTCTCCAGCACCAAGCT CCTGTGTCTAGCTGAAAACCTCTTGGAAGGGTCTGAAAACCTCAAACCTTTCTTCATT[T/C]T TGCAAGAAAAGAAGGCCAAGTCTTGCAAGTTCCCATCAGATTGGACTCTTCTGATCAAGC CTTTGTTTTGCCTTTCAAGCTCACCAATTGAGGGAGAGATTGGAGAAGAACCACATGCAG AGGACCCAGTTTGGTGGAAGTAAGTTTGTGATAATGTTGTGGGACAGTGGTTGGTGGA
>Gm03_4 0211426	TGGCGTACCAGCCGGTGTGCGAGAAGACGCCGTTGACGTTTTCCAACGGAGGACCGAGG CCGGCGTTGGTAGTGAACCTTGACATGTTGGTCCAAAGGGATAAGCTTCTGCAATGCTTC AACATGTCTCGTTGAAGCGCGAGGGAAGGTCTGTGGACGAAGATGTAGCGTCTCCGCA T[G/C]GGTCTGCTTGTGTTGTCGGCGGTTGCGAGGGCACGGTCTGAAGGGGAAAGTGGGTT CCTTCGGTTGCGAAACTCGCTCAGGGAAGCCGAGGTTTTTACGAGGTTGTTGGTGTGTT GTTGGGATTGGAAGTGATGGGGACTGCTGGGTCAGGGGCGGTGGGTTGGTGGCGGAG GACGGCGA
>Gm03_4 0371846	CAGAGCTCTTGAGCAAGAAGTGGTCTCCCTTACCGCATAGATCCATGCTTGGCATCCTT CATGCAGTAGCATTCTTTACTACCACTGAAGTGAGCCAAATGGTTTAGAATAAATTTTTTG GTTAGAGGAAGTGATATAATATTGGACATTTCTTCTACATCTTTGTTTCATACTTTCA[G/A]C AAATGTCAATTTAGAATTCAAAGCCAACCTTAACTGCTGGAAGGCTACCCCTTTTCGTT GATTTGTCACACACTTGCCTCACTATGGTGGTCTATATATACAAGTCCCAAGATACCATT CATTTTCACAAATATTGATGTCTTCAATAAGCATGTGGTCTCGTGATTTAACGTTGA
>Gm03_4 0459321	ATTTTCAATTTGTTATATGATTTGTTGACTTGTGTTTTGTTTATTTACTTATTTCACTTTTCA TGGACTTTGCAGATTGTTTCTGACTCAATGAGAGCTACAAATAAGAAAATAAATGGAACAC CCATCAAGATGCTCATAGACCAAGAAATGTCCAAAGAAGTTGTTTCCAAGCACA[C/G]CCC ACCACCAATGTAGTTGCAAAATTAATGGGGCTTGAAGCCCTCCACAGGGTGAGCTATC TGTGGAGAGAAGCCATAGAGGAGATTATTCTCAACATATGTGTGGTCATTACGGGACACC ATTCAACCACTGGAACCTGGAAGATAGGTTTATGGACAAGGAAATGCTTCATGAAGT
>Gm03_4 0462434	TGGATCCATCATTGAGGGACAAATACGCCAATCTCAATGACAAGGAGCCTCAGCAGCAAC TCCACGAGGCCAAGCGAAGGCAGAGGAGGTCCAATCAGAAGCTTGATTGCGATTGTGTG AATGTCTCACTAATAGAAATTAAGTTATGGATCAGAGAAGAATTACTTGATGGGTAGT[C /T]GGTTGTGCAGTGGGAGCCACAGCAGGGTCCAAGTCCCAGAAGCTGCATCTCCCCAT TGGTGGACCTCATTGTGGCACAGATGAAGGAGTTAATATCTAGTGCTATGAGTTCTGTTTG GGTGGTGATTGTGGGGACAGTAACAGCCTGGTGGTAGAGAGTGTGTGAGAAAAGAGG TTG
>Gm03_4 0462640	GTCCAAGTCCCAGAAGCTGCATCTCCCCATTGGTGGACCTCATTGTGGCACAGATGAAG GAGTTAATATCTAGTGCTATGAGTTCTGTTTGGGTGGTGGATTGTGGGGACAGTAACAGC CTGGTGGTAGAGAGTGTGTCAGAAAAGAGGTTGTGGGCAAAGGGTGGGTTGAGCTTAT[ G/T]AGATTGGAGATGGATATTTTGGTGAAGGAAGTAGAGGGGAAGCTGCTAGAAGAACTT GTGGAGGATGCGGTGGTTGATTTGACAGGCAGGGCCTGAATGGTTGAAAACCTGACAT GGCTTTCCGCTTTGGCTTTGCTTTTTTATAATATATATTGTAACCTCTACTATTGTTGATT

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>Gm03_4 0585266	TAAAGTGCATAACAATTGAGTATACCATCTAATAATGAAAATAACGCAAATAACATAATGCT TAAAGCAACGTTTCATATCAGAAGGGAAAAATAGTTTGAATCCCATCCAACAAAAATCATGC GATTAATAATCCAATTAATAAGCATAAAGTAAATAAGTTTTTAAACAGGTATCT[T/G]GAG GAAAGCAACTACATTTAGTCCACTTCCTCAATCTTGGGGCCAGCACCACCTCCACCAGAA GGAGGAGCAGCATACTCATCCTCTGCTGCACCAGCACCACCCACGTCTGGACCAGCACC ACCTTGGTACATCTTGGCAATGATTGGATTGCAGATGCTTTCCAATTCCTTCATTTTG
>Gm03_4 0585499	TCCACCAGAAGGAGGAGCAGCATACTCATCCTCTGCTGCACCAGCACCACCCACGTCTG GACCAGCACCACCTTGGTACATCTTGGCAATGATTGGATTGCAGATGCTTTCCAATTCCTT CATTTTGTCTCAAACCTCATCTGCTTCTGCAAGCTGGTTGCTGTCTAACCACTGGATTG[A/ C]TTTGCTCAATTGCATCCTCAATCTTCTTCTTGTGTCAGCCGGTCAAGTTTCTCACCATCTT GTCATCCTTCACGGTGTTCCTCATGTTGTATGCATAGTTTTCCAAAGCGTTTTTGGCCTCA ACCTTCTTCTTGTGCTCTTCATCCTCAGACTTGTACTTCTCAGCCTCTTGAACCATCTT
>Gm03_4 0586108	GTCAGAGTATGTTGAGAAAACCTTGTTCTTCTTTGTTGGAATTGTAGTGTTCCTAGGGATC AGGACAGTCATCACACCACCGGCAGTCTCCAAACCAAGAGACAGAGGGGTGACATCCAG GAGGAGAAGATCCTGAACCTTCTCATTGCCCTCACCACCTAAGATTGCAGCCTGAACAG[G/ T]JCGCACCATATGCAACAGCCTCATCGGGATTAATGCTCTTGCAAAGCTCTTTTCCATTAA AGAAGTCCTGCAGCAGTTGTTGAACCTTGGGAATTCTGGTAGAACCACCAACAAGGACAA CATCATCAACACTTCTTTGTCCATCTTAGCATCCCTCAAACATTTCTCCACCGGCTCCAT
>Gm03_4 0587775	AATAAGTATAAAAAAATACATAAACAGAACATCGAATGAAAAGATAAAATAATTTTGT AAGAGAGGTTACCGAAGACGGTGTGATGGGGTTCATGGCGACTTGATTCTTGGCCGCAT CACCGATGAGACGCTCGGTGTCAGTGAATCCGACGTAAGACGGCGTGGTTCTGTTCC[T/C ]TTGGTCTGTTGGCGATGATTTCAACGCGGTCTGTTGCCACACACCGACGCAAGAGTAGG TGGTTCCGAGATCGATTCCGATAGCAGGACCCTCTCCTTTTCCGGCCATGGTTTCGGCGT GCGGCGTAGAGAAAGAAGAGATCTAAGAAAGAGGAAGAAGAAGATGATAACCACAGCGA TA
>Gm03_4 0600203	CTACGAAAACGCACAATTGGTATTGGCAGGGTAAATTGGTGAAGTTGAACAAAATGAAGC TAAGAAATCACTGATGGAAGCGGAAATTGGGGAACGATCTGATCCGTGAGCGAAAGGAA GGAAATCAGAGACGAAATTGAGAGTGGTGTGGTTACCTTTGGACGCGGTGGCAGTGGGC[ G/T]TGACTTTTCCCTCTGATGAGTGTGTTGTCGTGCTTCTAGGTCTCGCTGCGGTGTTT CCCCACAGTCCCTCCAACACAAATAAGAAGCAGAATAGTGTCTTACTTAGTTGACCATTTTG CCCATCTCTATTTCCCAAATTTAGATAATAAGGTTTAAATACCATTTTATTCTTTTAAATT
>Gm03_4 0600256	ATGAAGCTAAGAAATCACTGATGGAAGCGGAAATTGGGGAACGATCTGATCCGTGAGCG AAAGGAAGGAAATCAGAGACGAAATTGAGAGTGGTGTGGTTACCTTTGGACGCGGTGGC AGTGGGCGTGACTTTTCCCTCTGATGAGTGTGTTGTCGTGCTTCTAGGTCTCGCTGCG[ T/C]TGTTCCCCACAGTCCCTCCAACACAAATAAGAAGCAGAATAGTGTCTTACTTAGTTGA CCATTTTGCCCATCTCTATTTCCCAAATTTAGATAATAAGGTTTAAATACCATTTTATTCTTT TAAATTTGGCTGATTGTTTTTTAGTATTTAAAAAATATATTTTTATTAGTCTTCAA
>Gm03_4 0628097	ATTTCGAAATAGAATGACGAGGAGGATTGAGGACTTCGTGTCCAGTTGTTCTTCCCTT GTACTTTGCTGCCAGTGGTTTGAAGAACTGACGTGACTAAGTTACGAAGCGTGGTGGATTG GGGGCTTCTTTGCTGGTTACGTCCACCGCGAGCGTGGGGAAGATTTTGGGAACGTTTG[ A/G]CGTGGCGATGATGTGCATGGTGTGGAATAAAATAAAATGAAGGAAAATAAATATG AAGAAGATGCACAGTCTTGAATTAATAACAAAATAGCAGTAGCAAATTAATTTAATTGAC AACACATAAATAATGCATTATATCACACAATGAGCACAAGGAAAAATTAATTAAGGGAA
>Gm03_4 0676583	CCTTTCTAAAATTTTCAGTATTTTCTCATTGAGCAAAAAGCATACATTGCTTACAACTATACC TTCAAATTTACTAACAGCCAAGTAATATAAATCAGATAACTTACAGGCAGAGATACAGAG CGTGAGTAAGTTTGATGAGCAGAATGCTCATAGAGTGGTTCTGTGACTGAAGTATC[G/A]G CCTCCTTATCAGACGAGCTTGAAGGCAACATGGTATTAGGAATTGGAAAGGTTGCAGATT TTTTCAAGCACGCCATATTTACTTTTGTGCTGTTGATCTTTCATTGCTTTACATATTTTTCCT CATTCATTATAGATGTCACTCCCTCAGAAGCAAGTAAAGATTTCTGAGCATCTT
>Gm03_4 0676856	TGATCTTTTATTGCTTTACATATTTTTTCTCATTACATTATAGATGTCACTCCCTCAGAA GCAAGTAAAGATTTCTGAGCATCTTGAATGTGTAATGACTCACTAAAAGTCCTTCCAAGAT TACTTATGAAGTTGTCGTATGCACAACATTGGTATTAGAGCACGTGTCCATCAA[C/T]TGC ATCAGGTACTAAGAGGCAACTAAAAAACAACAATCAGATCAATGAGAATAAATAACACAA TTATCACTAATCTACATAATAAGCATTAATAAATTAACATTCCACAGTTATGTTTCATTATT AATACATTGTGGACCAAAATTTAAAAAAGGTTTTAAATGATGACAGATG
>Gm03_4	CTTCATACATTACTAACATAAGTCGGAAAGCAAATGATACCTAAGCTTGAAAGTTGAACAA

0678154	GATTACCAAAAAATATACCTTATAAAACGACGTTTGGAGCTGCAACTACAACACTATCAA AACAGGAGCAGAGAATAGCAACCAACACAGAAAGAAACAACAATAGCAAACCCTCGA[C/T] ACCGATCCGCTGACGAATTAATCACAGCTCGCTCAGAGCAGAAACCAACAACGACGAC CAAAAGAGAGCGAGAAAAAGAAAGAGAAGGTGGAAGAAAACGAAAGACCTAGTGTTAACTA AGATTACCACGCCGATCACTTTTTTCAAGATTTTTTCAGACGATGAAAAGAGGATCCACCA
>Gm03_4 0683015	CCACGCTCTTCTTTGTTCTCTTCTTCTTCTCTCGCTTCCTCCTTCCTTCGATATCTTCTTCT CGGCCTTGGGCTTCTTCTCCGCCGAGCCTTCTCTGCGGGCTTTTTCTCCGCGGGCTTCT TCTCGGCCCTTTGGGGGAGCCATTGGTTGGTGAAGAGGGAAAGGAGAGGGTTCGGTTT[A/G] TTATGATGGGAGAAAAAACTGAATTAGATAGAACGAGTGAAGGAACAGAAGAATGTGTG TGTATATATAGGAAGAGAGAAAGCTTTTGATTGGAAGGGGTGAGTCACGCGGATCGGTG ACGTGGCGTCTTGGATTGACATTGCGTTTTTTCAGCGCCCAAGGATTGCTTCCGTCTTTT
>Gm03_4 0685721	ATCTACCATCAAAACAGCTAGCCTTAATGAAAAGGCCAAAAGTAAAAAATTGCTATAAGATT AAAAAAGTGAAGATAAATGTATTATTGTAGAGAAATCTAGGTCTCGCATCCTTTACA TGTAATTTAAAGCTCAAACAAATTCCTCGATGCGGGCTTCTGAGATTTAGACTCTGA[C/A]T AGCAGGGGAAATCAATCTCTGCTGAACACGGGAAATGAGATGTTGGCTATAACCTCTGCT CGCAACACTTTGAACCTGAATTTCCGATGAAGAATCATGTCTTATCTAGTACCTGTCAGGC TGTCACAGCAGGGGAAAACATGAGACAAGTCAAACGGGTAATAAAAAATAAAAAATGT
>Gm03_4 0785291	GGCCTTACCCATGGATTCTCTTCCCAAATTGCGCCACCCTTCTCCATTTCCCACTCCCTTA TCCTCTTCCCTCTCCTTCCGCTCTTTACCGCCACCCCATTTTCTTCGTGCTGTTTAGGT TACCCTCCATCAGAGCCTCCTCCTCACCCCTCCCAAAACGACCCCGCTTTTCGCACCC[C/T] AAGAAACAAGCCCTCCCTCCCCCCCCCTCCTCAAACCCTAACCTCCTTCTCTCCCCCT CCTCGAAACCACCTGCATCGTCATCGCCGCCGCCGCTTCTTCTTCATGCGCTTCCACCA CACGCCCGCGTCATCGCCGCTCCGCTCACTTCCCCCGCCGCGGAAACCGACACCGCCT T
>Gm03_4 0785299	CCATGGATTCTTCCCAAATTGCGCCACCCTTCTCCATTTCCCACTCCCTTATCCTCTTC CCTCTCCTTCCGCTCTTTACCGCCACCCCATTTTCTTCGTGCTGTTTAGGTACCCTCC ATCAGAGCCTCCTCCTCACCCCTCCCAAAACGACCCCGCTTTTCGCACCCCAAGAAAC[T/C] AGCCCTCCCTCCCCCCCCCTCCTCAAACCCTAACCTCCTTCTCTCCCCCTCCTCGAAA CCACCTGCATCGTCATCGCCGCCGCCGCTTCTTCTTCATGCGCTTCCACCACACGCCCG CCGTCTCGCCGCTCCGCTCACTTCCCCCGCCGCGGAAACCGACACCGCCTTACCCAG G
>Gm03_4 0823593	TACATCTTAATTGAAAAATCTATCATCGACGAAGGAATCTACCGCCTCGGCGTGGAGAAA CTTAGCTCCTCGCGCGCTAACAAGATGGATTGGGAAGTTCTTGATTTGAAGATCAAGAGT TGGTTAGAGGCGACAAGGATTTCCGTTAGAACACTCTTTAACGGAGAGAGAATACTCTG[A /C]GACCACGTCTTCAGTTACTCAGATTCCGTGAGAGAATCATGCTTCGCCGAGATTTCAAG AGACGGTGCCGCTCTCCTTTTCGGATTCCCTGAACTCGTTGCCAAACAAAGAAATCTTC GCCGGAGAAGCTGTTCCGCGTGTCTGATATGCATGCTGTGGCTTCCGAAGTGTGCCGG AG
>Gm03_4 0874888	CCTGGGGTGACCGTACAATATCTGAGAACCCTTTTCTCTGTTGCAACTTCTAGACCTCAG AACTCAGCATCCGATTCCGGCCACTTATAGAGGTTACGTAGGAAGCCCTACCCGAACA CGTGGGTACGTGCGTCTGTTGATGCAGTTTGATATGCATACTGAGCTCATCTTTTTT[C/T] CCCCTCAGTTTCTCAGGTGAAACTGTGATAGCTAGTGGGAGTTTGTGTAGAAGGGTATAA ATTGAGAATATTAAAAAAAGAAAAAAGAAAAAGGAGAAAAAACTTAGATTGCAA GAAAAGGGTGTGGGAAAGGAAAGAAAGGACTAATTGAATGTATTTATAGTGGGCATTATT
>Gm03_4 0889026	CTCCTAGGGAGGAAGTATACCAAGCAATTGAAGACTGCAGAGATGCTGGAATTCGTGTTA TGGTTATTACTGGAGACAACAAGAACCCGCTGAAGCTATATGCCGTGAAATAGGTGTAT TTTCTCCAGATGAAGACATTAGTTCTAAAAGTTTAAACAGGTAGAGATTTTATGGAAGT[G/A ]ATGATAAAAAGGCATATCTGAGACAGCATGGCGGGCTTTTGTTCGTTTCAAGGGCTGAACCTA GGCACAAGCAAGAAATTGTGAGGCTGCTCAAAGAGGAAGGGGAAGTGGTGGCCATGACT GGGGATGGTGTTAATGATGCTCCTGCCTTGAAGCTTGCAGACATTGGTATTGCAATGGGC A
>Gm03_4 1007691	CAAAAAAGAAGCCCACTTAGAAGGACTCTTTAAATATATTTATATTCAACGTATAAAACCT TTGATGCTGTTCTGCCAAACCCCAACCCCTAGTTTTGTGAGTAGCACACTTCACCCATGGC AGTAGCAGCAGCTCGCCCTATCGTCTCCGTCCAAACCCTAGAGGGCGATGCAACCC[C/G] TACCGTTCTCTTCCCGACGTCATGAAAGCCTCCATTTCGGCCTGACATCGTCAATTTCTGC CACTCCAACATTTCTCGCAACAGCCGCCAACCCCTACGCCGTCAGCAAGCGCGCGGCCA CCAAACCTCCGCCGAGTCTGGGGCACCGGCCGCGCGCTCTCGCGTATCCCGCGAGTC

	CC
>Gm03_4 1007937	CAACATTTCTCGCAACAGCCGCCAACCCCTACGCCGTCAGCAAGCGCGCCGCCACCAAA CCTCCGCCGAGTCTTGGGGCACCGGCCGCGCCGTCTCGCGTATCCCGCGAGTCCCCGG CGGTGGTACCCACCGCGCCGCCAGGGCACCTTCGAAACATGTGCCGTGGCGGCCGC ATGT[T/C]TGCTCCCAACCAAGATCTGGCGCCGCTGGCACCGCAAGATCAACGTTCAACAG AAGCGCCACGCCGTCTGTCTCCGCCATCGCCGCCTCCGCAATCCCCTCGCTCGTCCAGGC TCGCGGCCACCGCATCGAGTCCGTCCCGAACTCCCCCTCGTCTGTCAGCGACACTGTCTG AAAGCGTCGA
>Gm03_4 1008442	TACGGAACAGAAGGTGCTAAGGCCATTAAAGCCTTCAGGAACATTCCCGGTGTAGAGGTG GCAAACGTTGATAGGCTCAACCTTCTGAAGCTGGCACCCGGTGGCCACCTTGGAAGGTT CATTATTTGGACCAAGTCCGCATTTGAGAAATTGGACTCTATCTATGGATCCTTCGATAA[G /C]GCCTCCGAGAAGAAAAATGGCTACTTGCTTCCCAGGCCCAAGATGATCAACTCTGACT TGGCCCGTATCATTAACTCCGACGAGGTCCAGTCCGTGGTCAGGCCCATTAAGAAAGATG TTAAGAGAGCACCTCTCAAGAAGAACCCTCTCACGAACCTCAATGCGATGTTGAAGCTCA AT
>Gm03_4 1008459	TAAGGCCATTAAAGCCTTCAGGAACATTCCCGGTGTAGAGGTGGCAAACGTTGATAGGCT CAACCTTCTGAAGCTGGCACCCGGTGGCCACCTTGGAAGGTTCAATTATTTGGACCAAGTC CGCATTTGAGAAATTGGACTCTATCTATGGATCCTTCGATAAGGCCTCCGAGAAGAAAA[T/ C]TGGCTACTTGCTTCCCAGGCCCAAGATGATCAACTCTGACTTGGCCCGTATCATTAACT CCGACGAGGTCCAGTCCGTGGTCAGGCCCATTAAGAAAGATGTTAAGAGAGCACCTCTC AAGAAGAACCCTCTCACGAACCTCAATGCGATGTTGAAGCTCAATCCCTATGCTAAGACC GC
>Gm03_4 1171468	TTCCGCAAAGAACCCTCACCCCTTTATTGAAAGGCCAGTACCTCTCAATGTGGGTGTCATA GTACAATAATGATCCCAGGGAATATATTGACAATCAATATTTGGCTTCTCCTGCCACAATT CTGAAGATATAGGAAGATCAGTTGAACTCCAGCATTAAGTATCTACTAGTACTGCC[C/T] CCCTCATAGACTGAAGGAATTCAGGGCTCTTACGCTTGATGCCCTTGTCCTACTTCCC ATTGTAAGATCAATAAAACCACCATCAACATGAAATGTTTGTCTGTCAATAATTGGAGGAAG AAGGTTGGTGTGCATAATTTTCCAACGGTATAAACAATGATGGTCCAGCACTAGACAAA
>Gm03_4 1185505	CTGGCATGCAAATGTTCAAAATTAATGGGGTTATACCTTTGTCCAATGCATGGTACAATGT ATTATTGCAGGTAAAGGGGCTTTCATCACAAGAATTTGCTGCCCGTAATGATTTGGCTCTT GCATTGCCGATAGGATTCAAGCTATCCCAGATGGGACCCCTGCAGCATCCAAACAA[C/T] GTGGAAGTTGGGCAGCTTCAGCCTCACGTCTGGAATTAATTTGATTCAAGGTAATTGAAA TTATCCATGTCTCATAGAAGTTTATCAATGGGAGACTTTTTAATTGATGGTCATCCATTCT TTTTGTGTATCTTGTCAATTCCAGATGGGAAATTCGATGATGAATACTTCCAACAAA
>Gm03_4 1223681	TATCAGCTATTTGTGAGGCTTTGAGGAGTTTGATGAAGGTTGTGAATTGTCCTGCTTTCTC AAGCACTTGGGTGATGTTGGTGGGTCCCGCTGGTGCCGGTGCTACTTGGGCTGAAATTG TTTGATGAAGATGTTGAGGAAGAGGAAAGGGAGGAAATTGAATAGCTTTGTCAATTGGT[A /G]TTCCTTGTGAGTGAGGAAATTAAGAAGAAAGACTAAGATGTTTGTACAAGTGGGAAG GAAGAGATGTGGTGTGTAATATATATTAAGAAGAGGGTGCATGGAATTTGATTGTTTAA AAGGTTAGGTGAAAGGTTGCAATTAGTATAAGGTAGCAAGAGGCCCTCATGCAATCAATT T
>Gm03_4 1228284	TAGGCCTTGGTTTGAGTTGTTGAGCTGTGAATTGATTCTATCAGCTATTTGTGAAGCTTTG AGGAGTTTAAATGAAGGTTGTGAATTGTCCTGCTTTCTCTAGAACTTGGGTGATGTTGGTGG GGCCCGCTGGTGCTGCTGGTGTGGGGCTGAAATTGTTTGATGAAAAGGAAAAGGA[G/A ]AAAATGGATTAGCTTTGCCATTTGTGTGAGTGAGGAAGCTAAGAAGACTAAGATTTTTGTC ACAAGTGGAAGAAGGGGATGTGGTGTGTGTATTTATATTATGGTGTTAAGGTTTGAGG GTGAGTGGAATTAGATTGGTATAAAAGTTAGGTGAAAGGTTGCAATTTGTATAAGG
>Gm03_4 1228321	CTATCAGCTATTTGTGAAGCTTTGAGGAGTTTAAATGAAGGTTGTGAATTGTCCTGCTTTCT CTAGAACTTGGGTGATGTTGGTGGGGCCCGCTGGTGCTGCTGGTGTGGGCTGAAATG GTTTGGATGAAAAGGAAAAGGAGAAAATGGATTAGCTTTGCCATTTGTGTGAGTGAGGA[A /T]GCTAAGAAGACTAAGATTTTTGTACAAAGTGAAGAAGGGGATGTGGTGTGTGTATTT ATATTATATGGTGTTAAGGTTTGAGGGTGAGTGGAATTAGATTGGTATAAAAAGTTAGGT GAAAGGTTGCAATTTGTATAAGGTGGCAAGAGGCCCTCATGCAATCAATTTTTCTTTCTA
>Gm03_4 1234338	GATGATTTAGTTCAAGTTTGAGCATGGAAGATCGCAAGGTTGTGGCATGGGGCGAGGG ACATAACTCATGGGTAAAGTTGCCTGACAATTTTTAGTTGTGTGCTTGTAGCAAACCTTGCT ACCGTAGGGTTAATGTAGTGTGTATCAAATGCCATGATCAGGTGAGTGGAGTTGCTT[T/ C]TGATTCATATTGGACATCACCAAATCAAATGACAATGGAGAGACTGTTACATATCGTTT

	TGGTTCTGTTGGTCAGGTGCTCAGAAAAATGTTTCATTATAATTTTTACTTTTTCTTTGTGT AGCTTTTCATTCTAAGTAAAACTCATTGGTATACCTAACATGGATCTGTGGTAGCTTT
>Gm03_4 1274006	AATATACACTGACAAAGTGATCCTCAGACCTCTATTGAAACGGTGATACATATAGGCATGC CTATGAATTCTTAACTGTTTATTTTTAAAACTCAGAAATTATTTTTAATACATGGCAGCAAG GTACCACGTATTATAGCCAACAAAAAAGCAAGAATCTATCAGCGCCAGGATCTAG[G/C]AT CAACAATAAACACAATATAAATGAGATATACTGTTCTGCATAACTTAAGAGTTCCCCAGCA CCAATATTTCTCTAATTCTTCACACTTGTTACATGCTTGAGGAGCATGGAGACAACATCCC CTGATATCCTCTCAGCCTCAGTTTTCAGGTGATGAATCTTTCTTCTGTCTCTTGC
>Gm03_4 1275788	GAAAAGGAAATCGTGAGAAACGGGAGCGAAAAACGAGTTGATTATAGTAAAGAAAGCGCGT GAAATCCGAAAAGGGCTGCATTGGCGTGAAAATGGTAGATCTAAAAGTAAAGGAGAAAAAC CGAAATAAAGAGATTGGAAGGTACCGTAGCAATCGGGTCTGTGTTAGATTGAGAGAAAAAC[ C/T]AGACACAGTGTCTCTAACGAAAACCTGGAGGACTGCTTCGCTGATTTGCTTTTCGTCTT CGCCCAATCAAATCAACACAACCCACCTATTTCTTTAATAATAAATAAATCACACATATATA ATTTAATTTTACTCACAACCATATTCTTTTCGCAAGTGTGGACATAAAATAATATGACA
>Gm03_4 1982791	TGGATATGAGTGGAACAAATATAATCAGACTCACTATGATCATGACAACCCACCTCCAAAG ATAGTCCAAGGGTACAAATTTAATATTTTCTACCCTGATCTTGTAGACAAGACAAAAAGCCC CAACCTACACCATTTGAGAAGGATGGCAGCAATGGGGAGACTTGCATTATAAGATTCC[G/A] CGCAGGGCCACCATATGAAGACATAGTAAGTCTATTATCCTTTAATTTTATGAAGATTTTT AATTTTCAATCAATTTACTTATTTCTTATTTTTTCCATGACAGGCTTCCGCATTGTAAACA AAGAATGGGAGTATTCTCATAAGAAAGGTTTTAAGTGTGCATTTGAACGTGGAAT
>Gm03_4 2074261	AGGCCCACTACTCTGCCGTTTGGGGAGCATTCCACGCCGCGCCACGCGCGGAGCAGG CGTCGGCGCCCGTCCAGTTGGTGAGGAGTTGCCGTGGGTGTCCGTTTGGAGACGGAA TTCGGTGAGAGCGAGTGTATCATTTTGCCTGCAGCCTCTGCAGCAACACAGAGGATGCA CAA[C/T]CACAAGGCCAGTGCCAGGAATATGGGACAATGACAACTTTTTTCATGTTATGG TCTGGGAAGGGAGGGAAACGAGTGTGGTGGTGTATTAAATTTAAAGAAATTCAGATTTGG ACTGAGAAGGAAAATGTGTCATGATTGGAGAGAGTGTGTGATTGAGGAAAAGTGTGAAG CTGTTA
>Gm03_4 2224757	CGTATAAGAATCTCGAATCCGCAAGTTTTAGATCCATTGAAATGAGCAATAGTAACAGTGC CAACGAATGACGCAAGGGGTGACGCATGTGCTCTTAGTTGACAATTGGTGGGGTCCATCT AATCTAATGTTTGTCTTCTGTTGGTGTGAGGTCCGGTGAAAAGGTACCGGATTGTGT[A/T] GGTTTTCTCACCACCCAGGGGCGCAACTCTCGCAAGTGGCGGTTTCAAAAACATGACTTC ACTACAGTGTTATGATCATAGCATGGTAAAGTCAATCAGCTTCATAGACTGAACTTAAACT AAATAAACATTTGTCCGACGTGTCTTTTTTATATTTCAATTTGACTTGAAACTGCCGCC
>Gm03_4 2224778	CAAGTTTTAGATCCATTGAAATGAGCAATAGTAACAGTGCCAACGAATGACGCAAGGGGT GACGCATGTGCTCTTAGTTGACAATTGGTGGGGTCCATCTAATCTAATGTTTGTCTGT TGGTTTGAGGTCCGGTGAAAAGGTACCGGATTGTGTTGGTTTTCTCACCACCCAGGG[C/ T]CGCAACTCTCGCAAGTGGCGGTTTTCAAAAACATGACTTCACTACAGTGTATGATCATA GCATGGTAAAGTCAATCAGCTTCATAGACTGAACTTAAACTAAATAAACATTTGTCCGACG TGCTTTTTTATATTTCAATTTGACTTGAAACTGCCGCCGCCGCCGCCGCCGCCGCCGCC
>Gm03_4 2243466	GCTCAAGATCGGCCCAAACGAACCATCCGAGCTGGCTATCCATGAAAATGCCTACGGTCT TGCCCGATATGCCGTATATGCCAGGAGAATGGTCTGGTACCTATTGTAGAGCCAGAGAT CCTGGTGGATGGACCTCATGACATCAACAAGTGTGCTGAGGTGACCGAGCGCGTTCTTG[ G/A]AGCATGCTACAAGGCTCTAAATGATCACCATGTTCTGCTTGAGGGCACTCTGTTGAA GCCCAACATGGTCACCCCTGGTTCAGAGTCTAAGAAGGTCACCCCAGAGGTGATTGCTCA ATACACTGTTACAGCTTTCAGCGCAACTGTTCCCGCTGCTGTTCCGGCCATTGTCTTCTTG TC
>Gm03_4 2243699	TGAAGCCCAACATGGTCACCCCTGGTTCAGAGTCTAAGAAGGTCACCCCAGAGGTGATTG CTCAATACACTGTTACAGCTTTGCAGCGAACTGTTCCCGCTGCTGTTCCGGCCATTGTCTT CTTGCTGTTGGTGGCAGAGCGAGGAGGAGGCAACCCTCAACCTCAACGCCATGAACAAG[ G/A]CCCAGGGAAAGAAGCCGTGGTCCCTTTCTTTCTTTTGAAGGGCACTTCAGCAAA GCACTCTCAAGGCATGGGGTGGGAAAGATGAAAACATTAAGAAGGCTCAGGATGCTTTAT TTGCCAGGTGCAATGCAAACTCACATGCAACTTTGGGAACCTTACAAAGGTGATGCTACCC TTG
>Gm03_4 2672131	CCCCAAAACAAAAAAGACAATTGCAGATCAATACGCAGCGAATGTAGAAGAACATGGAA AATTAATAAATAAACTAAAAATTAATAAATTAAGTGGAAAGTGAGAGAAACACTAACTT CTTCAAATCCGGTGGTTGCCCTGATCTGCTCATTGTGAGGAGAAGAGATCACAGATG[C/T] CAACGAAGGAAGAAAATTTGGTTTCGAAGTGGCAAAACCTCAACTACGCTCACTCAAGAC



	CCTAACATCGGAGGCTATACTCTTTATCGTTTTAAAGCCAGGCCAGTATACGAAACGATTGCTCAAAGGCACATGGATCTGATAGCATTAGTAAACAGCCAGGACCAAAATATGGGAAAG
>Gm03_4 2701398	GGATTGGGACGAAATGGTTCACCTGTTGTGCATCATGAGCAGCTCCAAGTCCTTCACATTGTGAACAACAAATTTCTTGAACCATTAGGGAGATAGTGGCGGGTCTTCTTGTCTGAACCA TAACCAATGTTTGGCATCAAAGTGCATCCCTTGAACCTTCTCCTAACACGAGAATCTA[C/G] GCCCTTTGGCCTGCGCCAGTTTGTCTGTTGAAATAGTATAATAAAAAAATGAGGTGAGAAA ATAAGAACTGATTGTTTATACACACTAGGTTAATAATCTTATTTAGGATAATAATTTTCATATT CAATCTTAGCCGAAAGATGGTGATATTTAGGATAATAATCATAGACAATCGGCAAA
>Gm03_4 2701407	CGAAATGGTTCACCTGTTGTGCATCATGAGCAGCTCCAAGTCCTTCACATTGTGAACAACA AATTTCTTGAACCATTAGGGAGATAGTGGCGGGTCTTCTTGTCTGAACCATAACCAATGT TTGGCATCAAAGTGCATCCCTTGAACCTTCTCCTAACACGAGAATCTATGCCCTTTG[T/C]C CTGCGCCAGTTTGTCTGTTGAAATAGTATAATAAAAAAATGAGGTGAGAAAATAAGAACTG ATTGTTTATACACACTAGGTTAATAATCTTATTTAGGATAATAATTTTCATATTTCAATCTTAGC CGAAAGATGGTGATATTTAGGATAATAATCATAGACAATCGGCAAAATTATTGAAA
>Gm03_4 5026222	CCTTTGGCCCTTCATCATAAAAAGTGTGTAATTGCTTCCTCCTCTTTGCTTGTGTTGGAT TTGAAAATGTTGAAAACCTGAGAAAAAAGAAGATTTCTTTGCCTGTTTTTGCACCCATGG TTATGATTGTTGATTTCTTGGCTATGAAAGACTATAGTGTCACTTTTAGTGACTAG[G/A]TAT AGGAACTTGTAGTATTTGTGGTGTGGGTAGAACTTCGAGGCTATGGGCCTCTCAATTTA TAGTACAAGTTAGGCCACTTCTTTAACTGAGATTAAGGGTGGAGAATCTGATTCAAACCA GTGTTTATGGTCCAATTAAGCTGACATGGAACATTTGAAACTGTTCCATTTTCAT
>Gm03_4 5416367	GGTTGCTTGTTATAGATGTTGATGCTGCCCTACCGTTCAATATTCCATTATCCATAGGTGG ACTGGTTCCCCACTTACCTTCTGATTCACTAGGCTCTATCACTTTCACGGACCCATCTGTC AATCCAACAGCAAATTTGGTTGGGTCCAGTGGATGAGCCGCAATGGCAACAGGATAC[C/T] CAGATTGATTCCTGCAATATAGTATACCACAATATGGATCACCAAATGGAAGTAAGACTTC ACGATACTACATCCTTAAATAAATGCTCAATGTTTAAACAATAGAGAGTAAATTTGGGTCT GATCTGTTGGGCAGAAACTTACCCGCTTAAAGCTGCTGGAGGTGAAAAGTATGTGGA
>Gm03_4 5503654	CTAAATACACAAATAGAAAATGTTTTGAAAAGGAGCAGGGAGTGCAACATTATAAAGAGTG GCCACTCAATAGTCAATAAGGGGAATCAAAAAATCAAATGCTAGTTTTTTGTAATACACGT CAATGTAAAACATGGGAGCCACGATATGTTATCAGAACAGAAATATCATAAAAACTG[C/T]C ACCGACGCCTAGAGCAAACAGTTTAATTTGAGAAAGTGAACAATTGTAACAAAAAATAGAA CAAAGTTATATTATTTCTTAAACTCTAAATCCTCTGCTGTTCTCCTCTAGCCCTC TCGAACCTTCTTCCCTTGTCTCGTACATAGGGCTTGGTATGGCTGTGTGGCACACC
>Gm03_4 5503673	ATGTTTTGAAAAGGAGCAGGGAGTGCAACATTATAAAGAGTGGCCACTCAATAGTCAATA AGGGGAATCAAAAAATCAAATGCTAGTTTTTTGTAATACACGTCAATGTAAAACATGGGAG CCACGATATGTTATCAGAACAGAAATATCATAAAAACTGTCACCGACGCCTAGAGCAA[G/T] ]CAGTTTAATTTGAGAAAGTGAACAATTGTAACAAAAAATAGAACAAAGTTATATTATATTTT TTAAACTCTAAATCCTCTGCTGTTCTCCTCTAGCCCTCTCGAAGCTTTCTTCCCTTTG CTCGTACATAGGGCTTGGTATGGCTGTGTGGCACACCAGGAGCAGGACCAAAAGTGT
>Gm03_4 5516926	AGGAGGAATGAACAATACCAAAAAACAAACGCAGAATGGCTTATTAATTATAAATATACATT ACATGATAAGGAGTAAAGATACACCGCTACTCTACCCTTACACTTTCTACTTACGATTTGG CAACCACTACGCAGCTTACTTAAGTAAAGACTTAAAGAAGCTAAACCAAAAGCTACC[A/G] CGCATGAACCTTGGCAGAACTTCCTTGCAGACATTGAGACATTATACATGCACACAAAGACT AAACAAAAATGAATACAGCAAAGGCCAAAAAAGCTGCTACATCCTCAGCTTTCAGGTACG AGAAAATGCGTCACACATCGAAAGGAGAAGAAGGGCCTGATCTGTTCTGCCCTCTCTTC
>Gm03_4 5516951	AAACGCAGAATGGCTTATTAATTATAAATATACATTACATGATAAGGAGTAAAGATACACC GCTACTCTACCCTTACACTTTCTACTTACGATTTGGCAACCACTACGCAGCTTACTTAAGT AAAGACTTAAAGAAGCTAAACCAAAAGCTACCTCGCATGAACTTGGCAGAACTTCT[G/A] GCAGACATTGAGACATTATACATGCACACAAAGACTAAACAAAAATGAATACAGCAAAGGC CAAAAAAGCTGCTACATCCTCAGCTTTTACGGTACGAGAAAATGCGTCACACATCGAAAGG AGAAGAAGGGCCTGATCTGTTCTGCCCTCTCTTCTCGTACGCATCAGATCCCGACCTTC
>Gm04_4 053359	ATTTGTATGCTAATGTGTTTCTTGTATTGACATAGGATATCATTGCTCTTCCAAGGTTGTTT CGGTTTCTCCAGCGACCATTTGGCAAAATTGATTTCTGTACTTCGGGCTCCTAAATCCAAGG AAGGGTATGCTGCTATTGGTGGTGGCTCTCCTTTACGAAAAATTACAGATGACCAG[A/C]T GGAGTTTAAATTTTTTGGTTTTCCCATTTATCTGCTTTGTGGAGCTTTTATCTTTCCGCAACA TGAATCTTTTTTGGCCTTTTCTGATACACTCAATAAAAAATTCTCTGTATTGGTAATGTGCT GGAACATAATGAAGCTGGTTCGTTGTCCATGAGCAGGCACTTGCAATTAATGAGG
>Gm04_5	ATGCCTCAACACGTAAAAATGAAAAGAAAAAATAACACCCCAAAACAAATCTAAAAAA

032005	TCACGAGCAAGTGGTTAAATAATAAGATTTGAAGAGAAGAAAGAAGGGGAAAAGGCGTT ATTACTTCTAGGAGTGCGAGGAGCTCCTCCACTGACGAAGGAGGATCTACAAGCTTGTT[/ C]JACCGGCTTCGAGAAGTTGTTCTTCCAATTCTTTATCTGCTCCCGCCATTGGATTGGTG GTGCCGTCTCTGAGTCTCTCTTCTTTCTGATCTAACACGGAATCGCGCCACACACTCTCTT CTGTTTCACTCCATCGATGCCCTGAAAACCTCCAATCGAATCGATTGAGATGGAAAGAG
>Gm04_4 4787569	GGTGGGGATAGGCGAAGGAGGTGGCGACTGAGTAGTGAATAGGGTTTTTGATTGATTGA TAGCAATAATTGGTGTTAGCGAAGCTTGAGAATTGAGATTTATGGGTCGCAATCGCAGTC GCAGTCGCAGTGTGTACGCTCTCCTTCCAATTCGCCGAGAAGAAGGTACTCTCCATCCC[ G/A]TGTTCCTCATCGCCACTCGAGAACTTCAGGCGCCGCTCTCCTTCCACAAGCGCCG TCGAAGGCACAGAACCTCTTCTCCCCCTCTCTTCACGCAGCCCCACTCCCAAACCCAA GAAAGATCAAAACAAAAGGTAACCAATCATAACCATTCTTTATTTGCCTGGATTTTATCTT A
>Gm04_4 5061509	AGAAAGAGCTGTTCAACTTTGTGATGAGTCACTTGGATGCAGCCACACAAGATGAAGAAG GGTCTGCACCTGTTTTGTTTACCCATTTGGGAGCTACCTTTTTGTTTCATCAGATGGAGC AGTGTGTGTGGATTGTGCAACTTCTGTTATAAAGATGTTGCATTTGATGTTTGCATCA[C/T] TTTCCCCAGCTTGTAGTATTGAGGACCATCTAAAGTATGGAGACTATGTTAAGAATTTGTC ATTACAATTCTTGAACCAGAACAAATTCGTCAGGGGAATATTCCTGATTCAAGCTATGAA GCAGGACTTGAATTGTCGGTTCAATCTTCAGGTTTGGGTAACCAGGTAATGCACTAT
>Gm04_4 5090419	CCAACCCCCCTCCCCCAATAAAAAATAGGAAAAGAAATTAGCTAACCCTAATGACTGATGA TGTAACAGTTATGGAGGAACCGTTGGCAGCTTGAAGAACATTGCGACTAGAAACATGTC AATTGCTTTGCAAACCAAGGCATACAATCTGACTGTTTTTGCCTGTATTCTTACAAAT[A/T]C TGTTGCAAAATTGGTGAGTTACTGGAGTTTGATTGGTTTTACTATTTGCAGAGACATTGA ATTTGTACAGATTTGCATGTATAATAATACATCAGTTTAAACAAGAAATTTGACTGAGAAAA TCACACGCCCCGTCCCTCTTGCAACTACTTTCACATTATTTACGTTAGATACTTCC
>Gm04_4 5157974	GTGGATTTGTTTAGCGTGGAGAGGAAAGCTTTGTTGCAGAGTTTGCCTTTGACGATGTTT GGGTGTTTGAGATTGGGCGAACAGAAAGATGATGATGGCCATTTGTTAGACTTGTGTTGC GTTGCGTGCACTAGTTGACTACTTCCCTTTACATTCCCTTCTTCTTCTGTTAGCACC[A/ T]TAGGTGCATTCCTATCCATCCCTTAGTCAAAGCTTTTCTAACTCTAATAATTCCTTTTAC TAAACTTTTCTTCTAGACGCTTGCCACCTCATTCTCAGGGATCCTTCTCTGCCATTTTT TTTTTTTACTAAAATGGGTAAAAATTAATAATTATAATAACGTTGAACAAAA
>Gm04_4 5380790	GTTGTTTTCTTTCACAGCCAATGTGTCATCATCTTCAAAGAAACAATCAAGCAAAGATAGA ATGTCATTGTCAACTTGATTATTTGGAGCTGAAGGATGTGCTGAGCAAGAGGTCATCAATT CTGGTACACAAGATACAGAGCCAACCTTGATTGCTTGGTGCTGAAGGATGGTCTGAGA[A/G] TGAGGGCATCAATTCTGATGCAGATGACACAGAGCCAACCTTGACAGTTTGGTGCTGAAGG ATGTGCCGAGCAAGATGGCATGAATTCTGAAACACGTGAAACAGAGCCAACCTTGACTGTT TGGTGCTGAAGGATGTGCCGAGCAAGAGGGCATGAATTCTGAAACACATGAAACAGAGC C
>Gm04_4 5594453	AGGTAATAAAAAAGTTAAACAGGAAACTTTTTCGCTTTTTCAATAATTGGTTGGAGACAA GAACTATTTTTGTGGAAGACATCTTAGACATGTATAGTTTAGTTTATTTCAATTTTTCTTA ACAGGAAACAATCATTAGTAATTACTCTCATAAAATTGGTGATGCCAGCAACCAA[G/A]ATT TTGGTGTTGACTGCGATGAGCAGTAGTACATTAGGCCTCATTGAGGTTTATGACAATCATG GCCGTGCTTACCCTTGCAAAGGAAATGAACATGAGAAAATCTTAATCTTATAGATCCCTAG AAAACCTTAATAAACTCATCTACAAAATCATAGAAATGGACTCATAACTTCGTAA
>Gm05_1 77701	GTTACTGGGTGCACCTAGCATGTCCCGTGAATAAATGGGCCTCGGCCCGTTTATCTGACA AAAAAAAATAGAGAGAGAATTTTGCTCTTACCTATTTAATAAGTTCTACATAAACCCATTCC CTTTCCTCGAAAAATCACAGATATCTTGCTAGGGTTTTAGGAAAAGACACTGTCTCG[A/C]G AGTGAACACAGGTAATCTCAATTATTCAATCTTCTCTTTTCTTCTTCCAGTTTTCCCATGT TCTCTTCGTTTTTTGTTAATCTCACTTTTTTCCCTTTTCATTTCTTTTGATTGATTTATCT GTTTACTTTTTTTCGTGTAGATCAAATATATTGCTAGGTATGGTATTATACG
>Gm05_1 77702	TTACTGGGTGCACCTAGCATGTCCCGTGAATAAATGGGCCTCGGCCCGTTTATCTGACAA AAAAAAAATAGAGAGAGAATTTTGCTCTTACCTATTTAATAAGTTCTACATAAACCCATTCC TTTCTCGAAAAATCACAGATATCTTGCTAGGGTTTTAGGAAAAGACACTGTCTCGC[T/C]A GTGAACACAGGTAATCTCAATTATTCAATCTTCTTCTTTTCTTCTTCCAGTTTTCCCATGTT TCTCTTCGTTTTTTGTTAATCTCACTTTTTTCCCTTTTCATTTCTTTTGATTGATTTATCTG TTTACTTTTTTTCGTGTAGATCAAATATATTGCTAGGTATGGTATTATACGC
>Gm05_1 033474	ATGAAGACAAAATGCGGGAAAAGGGTGAGACCCAGTTATGTTCAAGGTTAAGAATTTTCA CGTCATTTTAAAAATATGGATTTATCCCAATTTGTGTTTGTGTAGTTCACACATTCTAGGAT

	GTCTTTTTGGCAGTCATTTGGGGCCTCCTCGAAGGAGAACAACCTGCAGAAGAAGAAG[G/A] ]GAGGGTTAAGCACCCGAAGCCTGAGGTAGGTTTTAGGTTGTTTGTGTGTGCGTGTGTT TATGTTTTCTTTCTTCCAAACTAAACTGGTGTAGCTGTCGATAACGGTTGAGTTACTGAT TTTTACTGCAGGACTCTGTTTATTATCATCCTACATTAAACCCTACAGGGGCTCCACC
>Gm05_1 128981	CTGCAATCCAACCAAAAAACAACCTTGTTCTTCTCCGTCCACGGAACCTCCTGATCGTTCA CCAAAACCATCACTCAGTCAATGCAAAATCAATAATCGCGTTTGACAGTTCCGATTAATTC AACAATTATGAGATTAAACAAGAGAAAAAAATGATGATCGATCGGAGAGACGGATC[C/T] AACCTCGCTTTCGCTCGCGCTCGCGGTGGCGCCCGGAGTTCTGAGGAGCGGCGTCTGTC GGCGGAGGCGTAGCCGGCGGCGACGGCGTCTTTGTTGTTGTTGGTGGTGGCGTCT AAAGGATGTTCTGACTGTGAGAGGTTGTTGATGCTGACGCTCTTCCTCATCGAATCGACG ACGA
>Gm05_1 128982	TGCAATCCAACCAAAAAACAACCTTGTTCTTCTCCGTCCACGGAACCTCCTGATCGTTCA CAAAACCATCACTCAGTCAATGCAAAATCAATAATCGCGTTTGACAGTTCCGATTAATTC ACAATTTATGAGATTAAACAAGAGAAAAAAATGATGATCGATCGGAGAGACGGATCG[A/T] ACCTCGCTTTCGCTCGCGCTCGCGGTGGCGCCCGGAGTTCTGAGGAGCGGCGTCTGCG GCGGAGGCGTAGCCGGCGGCGACGGCGTCTTTGTTGTTGTTGGTGGTGGTGGCGTCTAA AGGATGTTCTGACTGTGAGAGGTTGTTGATGCTGACGCTCTTCCTCATCGAATCGACGAC GAC
>Gm05_3 850942	TAACACAACAAAATCGGATTTTTAGTAGAAAATGGAAAAAGAGAAAAACTAAGGATGTT GAACATTGTGGGTATAGGCATCTGAACAGCAAAGGAAAAACAAGAGAGGAAGATAAAGTTA CCGGCGGTGACGAATCGGCGGTTGTATTGCATTCTCTTGAGACACGTCCGCGTGGCT[A/ T]CTTCTTCTTGCTTGTCTCGCCACTTTGGGAGTTTGTCTCTCACTTTCCCGCACGTGC GAGAGATCCGTGAACCTTACCTGCAATCCAAAGTAAATCATTAGAGTTTCGCACATGTTT GGAGAAACACCGAATAGAATTGAATACGCGTTGGAAAATAACTTACCCATTGTGATGAA
>Gm05_2 7712137	CGACCAGCGAGCGTATTATTAATTAACCCCAAAACAAAGAGCACTGTTTTAAAAA AAACCATTCTATCTTTTTCATGACACGAATGCGATAAACAGTGTTGAAGGATCTGTGCGTG AGTTCTTGGTGAAACAAAGGAATCGTAGATCCGTTTGTGTTGTAGTTTTTTTTTTT[A/G]GT CAGAGAAGAGAAGCGATGGGGTTGTGGGAATCTTTCTCAATTGGCTTCGCAGGTTAGTA GCTTTCAACTCTAACAAAATAACAATGTTGTTGCTCTATTTTTATTTTAAATTTATTTATTG GTTTTAACTCGATTAAATTGATTGTTGAGGAAAACGATTGTGGATTTTAAGGT
>Gm05_3 8251772	TGCTAATATTTCTTTGATACCTCTGTAAACCCAGATCTGTCTGGTAAGGTTGATTTGTGG CCAAATGAAGATGGTAAACTTGAGCAGCTTCACTCAGATGAAGCATATGAGATGATTCAAA ACCATTGAGTCTCATTCTCGGTAACAAAGCTGGAGATTTAACATCAGTAGCTGAAA[G/A] CAGCAAATTCAGGGTAGGGCAGGTCTATGCAGCATCAGTGATGTATGGTTATTTTCTTAAG CGAGTTGACCAAAGGTTCCAGTTGGAGAAGACAATGAAAGTTCTTCCAAATGCAACAGAA GAAGAGAATGGTGTTTCATCGAAATACGATGGACAATGCAAGACCCAGATTGAAACAGGA
>Gm05_3 8278658	AGGTTCCCTCCCTTCAGATCTGGCCTCCTGCGTCAACCTCCGCAACCTCTACATTACGCG CAACTTGCTCACCGGCCAAATTCCACCCTTTTTGTTTCATTTGCCGGACCTCGTTCGCTTG AACATGGGCTTCAACAACCTTCTCCGGCCCCCTTCCGTCCGCCTTCAACAACCTTACCC[T/C] ]TTTAAAACTTTGTTTCTCGAAAACAACCAGCTCTCCGGCCCAATCCCCGACTTGAACAA ACTCACCTCGACCAGTTCAACGTCTCCGACAACCTCCTCAACGGCTCTGTCCCTCTTAA GCTTCAGACATTCCCTCAGGACTCTTTTCTAGGTAACCTCCCTCTGTGGCCGACCACTCTC
>Gm05_3 8279366	GGAAGGGAACGCGAAGAAGTTGGTGTTTTTTGGGAATGCGGCGAGGGCGTTTGATTTGG AAGATTTGCTCAGGGCTTCGGCGGAGGTTTTGGGGAAAGGGACTTTCGGGACGGCGTAC AAGGCGGTGTTGGAGGCGGGGCCGGTGGTGGCCGTGAAGAGGTTGAAGGATGTGACAA TTT[C/T]CGAGAAGGAATTTAAGGAGAAGATTGAGGCGGTGGGAGCGATGGATCACGAGA GTTTGGTGCTCTCAGGGCTTACTATTTTCAGCAGGGATGAGAAGCTCCTTGCTATGATTA TATGCCCATGGGAAGCTTGCTGCCCCCTTTACATGGTCCGTCTTCTTTCTTTTTTTGTGCT TGCT
>Gm05_3 8280387	GGAGTGGAACCTCCCAAGATGGGTTCAATCCGTGGTTAGAGAAGAGTGGAACCTCTGAGGT CTTTGATCTTGAGCTCCTTAGGTATCAAAATGTAGAAGAGGAGATGGTTCAGTTGTTGCAA CTTGACGATTGATTGTGCAGCACAATACCCTGACAAGCGCCCTTCAATGTCTGAAGTGGT[A/ G]AGGAGCATACAAGAGTTGCGAAGGTCTAGCTTGAAAGAGGACCAGGACCAATCCAAC ATGACAATGATATACTATTATAGAAGAACATGTATCTTCTCCCTCGACTTTGCTTTGTGAGA GTCGGAGCCCATATTTTACTTGCTAAATCCTGTGCGTATGTTGCTTGCAATTCTGTTGGCT
>Gm05_3 8402327	CATAGGAAAGTACATAGAACAAGTCTACAAGGACCTTCCACCCACTCACCTGCTTTGTTG ACTACCATCTCAACCGATTGAAATCCTCTGCCCTCCTCGTCCGTCAAGAAATCCTACA

	AGCTTCCTGGATCTCATCATCTGCCTGCGCTCCAGCCCCAAAAGACCCGTGGTCTGTC[G/A] J]TCCAAGGCCAAGCCCCACCCCGTCTGGCCTGCTGCAGCCCTCACCGACAGCCCTGCTC TTCTATCCGCCAAAAGAGGCCCGGTCTCTCTAAAGATTGGTGTCAGCCCGCCAGGC CAAAGGGGTCTGCCACCCGGGACTGGGAGATCTAAGCTGCCAAGAGGCCTGGCCGTC CCCC
>Gm05_3 8432746	CATTAGTAAGACTTTCCGTCGCTGCTTGAGGTAAGTGAGAAACAAACATTTACTGGTCCA CCGGAATGTGAGAGATCATGTTATGGCTGCCACAAGGGTCTTAACAAGGGAGACTTC CAGAAGGCTTTTGACATTATTGTGCTCTTGATGTGTGGAATTTGTGAGAAATCGAG[T/C] CACCGTACTTGAATGCTGAAGGACAAAAACAAGGAGGAGGCATTGAGGACATACCTCTT TACCTTCTCTTCACTTTATGAATCGTTGAGCCTGGATCAACTACAAAAATTTTTGATCTCT CTGTTTTCGCACTCATAGCATTGTCAGTAGGATGATGATCAATGAGGAGCTCATGC
>Gm05_3 8433427	TGGGAGTGGCTATTCAAGGGGACGAGGAAGGGGTTCTGATGGAGGTTGAGGAGGACT GCTCAAAGGGGTTGAGCATTGAGGGGACCCAGGGAGATGGATCTACTCGCATGGTTAG CCTTAAGGGAGTTCGTGCTGAATAAGAGTTTTGATAATTTGGCTAACGCACTATAGCATT C[A/T]AGTTTTTATTCTTGATAGGCTTTAATTTCTTACGGATTTATCTTTGTTTATTGTTAC ATTGGAGCTTGCTTTTACCTATTTTACCTGCTTTTATTTGTTTGAATTGTACCC TTAAGTGATGCATATTGACCTTGCTCATTTACGTGAGCTATCAATGTCTTTCACACAA
>Gm05_3 8433580	AATTTGGCTAACGCACTATAGCATTCTAGTTTTTATTCTTGATAGGCTTTAATTTTCTTACGG ATTTTATCTTTGTTTATTGTTACATTGGAGCTTGCTTTTACCTATTTTACCTTG TTTTATTTTGTGTTGAATTGTACCCTTAAGTGATGCATATTGACCTTGCTCATTT[C/T]CGTGA GCTATCAATGCTTTCACACAACTTATCTTTGTTTTCTTTTGAATGAAAGTAATAATTT AAAGAAATTTTTCATAATTTTCTAATTACCTAATTTATCATTTGTATAAATATTAGGCTTAAAT ATTAATTTTAAACAAATAATAGTAAATTTTGGGCTCATTTGTTAGAAC
>Gm05_3 8589019	TGCTGAGGCGCTTGAAGTTATTCCCTATACTCTGGCTGAGAATGCTGGTTTGAACCCGAT TGCCATTGTTACCGAGCTGAGGAATCGTCATGCACAGGGTGAGATAAATGCTGGAATAAA TGTGAGGAAGGGTCAGATTACCAACATCTTGAGGAGAGAATGTGGTGCAGCCCCTTCTTG[ G/T]TAGCACAAAGTGCATCATGTTGGCGACAGAGTGTGTGCGGATGATTTTGAAGATTGA TGATATTGTAACGTGAGGTAGAGTTGATTGAACAATGTGGAATTTAGAGAGAACTGAAT CACCCATTATCTTCTTTTTTTTTTTTTTGACTTCGTGCTTTAATTTTATGTTGGTAGTTT
>Gm05_3 8596037	GTAGATAAAATAGAGGTAGAGGTGTGGTAGATTGAGAGCAACATTGAAATGCATGCCAA GACAGACTCGGAGGTAACAAGCCTGGATGCGTCGTCCAGCACAAGGTCTCCTCGGCGAG CAGTGTACTACGTTTACAGGCCCTTCCCACGATGGGAGAGAAACGACGACGCTCGTTGCCT [A/C]CACCCTGTTCTCAGCCCCATGGGTTCCCTCCTCACTCTCCTCCTCCAGCCG CTTCTCCGCTTACGCCACCCGAATAACCATAAATAAATCTTGGAAGGGCATCGACGT CATCGAAGAAGAGGGTCTTCTCCAATCCGAATTAGATCGCCAACATTCTCTCTCTCGTCGA TA
>Gm05_3 8696065	AATACCACAGTACAGTGC GTTCAGGAAAAATGAAAAACCAAGCCCCAGA ACTCGATTTCA AGTGGGTAACAGAGCAGTAGAGTTGATCCATTACACGTAGGTAGCTACTCTCCTCAGAA GAACCAATTTTTCATAAGCACCATGACGAATATCGAATATTTGAACTTAGTGGAATCAGG[A/ C]TCAAACATGAGGTGTATTCAAATGGTTGAGGAGGTTGAGGTTCTCATCCACAGCCA TGGCTGATGAAGAACTCTGCAATCCTGTTGCTGACCCACCAGCAGCAGAACTGGTAGATG AGGGCGTGTGAGTTAGAGCAAGCACTTCTGGCTCAGTAGGATGCAAGTCTCTGAGAAGC AC
>Gm05_3 8913395	AAACACAGAATATAAAAGAAACAGAAGGATTATATATCATATGATAGTTGAAACTTCTTCC AAGGGAAGTCTCTCTCTCTCTCTCTCCCTCTCTCTCTCTCTCTCCCTCTCTCTCCCTCCCT CTCTCTCTCTCTCTCTCTCTATATATATATATATATGTACATGTCTGTGTGTCTAT[T/A]GTAA GTTTCTAACATCTAAGGATAGCCCTGTCTCTGTCTGCCAATCCGAGGGGTGAGCGAGAGT GAAGGTGGTTGACATCTCTGAGAAGGGCACGCCTCATGTGTAGCTGAAGTGAAGGGTGC TTTATTTTCTCAATTTCTTCTTAATTTTGATCACTTGGCTGTGAACTGCAAAACCC
>Gm05_3 8913666	CACGCCTCATGTGTAGCTGAAGTGAAGGGTGCTTTATTTTCTCAATTTCTTCTTAAATTTG ATCACTTGGCTGTGAACTGCAAAACCCATGCAACCCTCTCGCGGAGGAGTAGGGGTCT GTTCTCCATCTTCTTCTGCAAACTCACAAACAGAGCCTTTTACAGGAACCAATATGT[C/T] TGAAATGTAATTTTGGCGTTAGGGATACCTATTCAAGGATATATATATATATATACCCGGAA GAAGGAAAGAAATATCAGATTTTGTAGAGAGGAAATAAGGAAAAAAATCTTTATTTATT ATTATTATTAATAAATGGAAGTTTATATGCGCTCCTTCCCGTCCACCTTCTCCATT
>Gm05_3 9082457	TACATAGAATTACTTAATATTTGATTTTTTTCATGAGAGTGATTATTATTTGCTGCAGCATGT GGTGCTAGTGGGGCTCTGTATACGCTGATTTGCTGTGTCTATTGGCTGTGGCTGCCTATAC

	TCTTGCTTCTACCGCCCCAAGATGAGACGACAGTATGGTCTAAAGGGAAATGGTTGT[C/T] CGGATTGCTTGATTGCTTCTGCGAGCCCTGCGCCCTCTGTCAAGAATATCGTGAGC TTCAACACCGTGGAATTTGACATGATTATTGGTATGCACTCATTATATATATTAACACTAC TCTATTTGCTAAATACCCAGCTATGCAAAGTTGGAACAAATTTCTTATTTTATAGA
>Gm05_3 9082469	CTTAATATTTGATTTTTTTCATGAGAGTGATTATTATTTGCTGCAGCATGTGGTGCTAGTGGG GCTCTGTATACGCTGATTTGCTGTGTCATTGGCTGTGGCTGCCTATACTCTTGCTTCTACC GCCCCAAGATGAGACGACAGTATGGTCTAAAGGGAAATGGTTGTTGCGATTGCTTG[C/T]T TCATTGCTTCTGCGAGCCCTGCGCCCTCTGTCAAGAATATCGTGAGCTTCAACACCGTGG ATTTGACATGATTATTGGTATGCACTCATTATATATATTAACACTACTCTATTTGCTAAAT ACCCAGCTATGCAAAGTTGGAACAAATTTCTTATTTTATAGAGAACTCCTTTT
>Gm05_3 9166124	CGAACCACAAGAGTAATTGATAGAGAGAAGGAAAAAAGATATTAGGAAATGAAGATGGAT GTGAAGAGGTTGAATACTGACCGACGACAGTGGAACCTTCAGAACCCATGTTTGAGGGTG AATTGGAATTGGATGGGGGTAGGGTTTTGGAATTGAAGAGAGAGATAGAAAGACACAAA[ C/T]TAAGAGAGAGAGAGAGAGAGAGAGAGGGGCGTGAATATGATCCTGAACAGACGATGA CTGTGACTCAACTCAAGTCTAACAGCTGTTTCGCAAAAGTAGAAGGTCTGTTACACTAACAC TAACACGTACCAATATATATAAATTAAATATAATAGCAGTACTACCAACAAAGTTTAATCAA A
>Gm05_3 9166125	GAACCACAAGAGTAATTGATAGAGAGAAGGAAAAAAGATATTAGGAAATGAAGATGGATG TGAAGAGGTTGAATACTGACCGACGACAGTGGAACCTTCAGAACCCATGTTTGAGGGTGA ATTGGAATTGGATGGGGGTAGGGTTTTGGAATTGAAGAGAGAGATAGAAAGACACAAAG[ T/A]AAGAGAGAGAGAGAGAGAGAGAGAGGGGGCGTGAATATGATCCTGAACAGACGATGAC TGTGACTCAACTCAAGTCTAACAGCTGTTTCGCAAAAGTAGAAGGTCTGTTACACTAACACT AACACGTACCAATATATATAAATTAAATATAATAGCAGTACTACCAACAAAGTTTAATCAAA C
>Gm06_2 168622	CTCCCTTCATCTACGGAGAATGTCTGATCAACAAAACCATGACTTCAAGCTGGGGCACCA TTTTCTGGGAGAGAAAAAGGGAAAAAAGGAGGAAAAAGAAGAAAAATAATACCCAGTTATAC ATTATTACTACTTTGTGTCAGGAAGCTTGTGTGTGTACGAAGGGGCTTTCATCTTTTTT[A/C ]CTTTTTTTTTTTGGTTAATTCACATGCTTCTTCTAGGACTTTTGAATGAGGGTTTTCTA TGGAATCATTACAATGATGGATGTCACGTAAAGCGGGCTTTAAAGACTGAAATCCCAT TGCTTTTATATGTTAATTAATGGTTTGATTTACTAAACGAGATTTTAAATTAAGC
>Gm06_5 456983	CATTCATTTGTGAACTTTTTTTATTCCAGTCTTTGAATGCTTATTGGTTTTATCAAAATGTGA AAGAAAATAATTTTTATCGACACTGAAAATACTGTAATTTGTCCATTTGCAGGGTTTAAAA CGGAAGTAGCTCTTGGATGCATCATGCAAAAGCAAAGGGAATAAGGAGCCGGG[T/C]AA ATTTGTTACTCTAGAAAAGTGCTGTGAAGATTGTTACAGTCAATATGTACATGAACCTTGCA TTTTTATTTTAATAACTAGTTTAATATCCATGTGGACCCACAAGTAAGTAAGGTTAAACATT TTTATGTTTATTTTAATAATGGTCTGTATCGGTTAAATATGCGTTCTTTACGT
>Gm06_1 3383322	TGAATGGTATGAATTTGAAGATAGTTCAATATCTATTGTCTTTTATATTTAATTTGCATCTA AATACCACCCTCAAGGTGCTTGTACCCCCACTATTTTTCGTCTTATTATTTCCCAATATCTA AGTTCTTCTTAGTTAAATATTGTTTATGATTTACAGGCATCCATTTTCTATAA[G/A]CTCCG CCCCTGGTGATGACTACCTTAGCGTTCATATCAGAACACTTGGTGATTGGACCCGGAGTC TCAAAGTCAAGTTCTCAGAGGTGAATTCAGGTTAACTTTTGTAGGATATTATGGTTTGGAT AATACTAATTTACATAGCAAACTCACAAATCCTTACATTATTACATCTGTTTA
>Gm06_1 5714602	GAAATTAACGATAATAATACCTATATTTCTTTATAGCGATCAAATGAATCACCCTCCAACCT TGCGTCTTATCTGCGGCCATAACAGAAAACGAAATCAGACACACCACCACCTCGCTCAGC CGGAACCCAGCGCCGCTTGTCTACCGTCTCGTCCGTCTTCCACCCCTCCGCCGCC[C/ T]CCGCCGACCTTGATCTCCGTTCTCTCCGCCGACCGCCGGAACCGCCTTTTCCGCCGT CCTCTGATCCTCCTCCGCCGCTGGTCTGTTGAACATCACCACCGCCGACGGCGGAT TTCCGGCGGTTCCCGGTGCTCCGCGCGCTTCTCCGGTGTGAAGTCTTGAGCTTGCCA ACGG
>Gm06_1 7838601	TGTTTAGTGTACCAACAATTTGCATGATGCCCAAAGAAAAACCCCTCAAATGAATTTTTGTG TAATTTGCAAAATCCTTATAAGAATCGTTTTAAAAAATAAAATTTGGTAGGCTTTATG AAATTTGTTGTAAGTACGCACGCACAAGAGTATCCCATGCTAACTGCAAAATA[A/T]GTT AAAACAAAAATAATCTGATAGTCGTATTTTATCAAAAAAATATCAAGGCAACATCGAAACA GAAATGTTAAACAGTTTCTGTACATCCATAAACACAGTTGATAACTTCCATGACTCCGA GGCTTAACAAGATCATACAAAAATATATAACTTTGCTTAAATGAAAAACAACAA
>Gm06_4 2683097	TTTTTATTACTACTATTATTTATTTTATTATAATATTTCTTATCCTTTAAATACATAGCTACTT AGCACATGATAGTTTGTTAATTTATATATCCAAAAGGCATTTTAGACCAAGTTCTATGATT

	TTGATTATAAGCTTTTGTAAACACAAAATAAGCTAATCCAAACATGCACCTTAC[A/T]GCTGC ATATTTAGGTCTACAGTGGGATCAGCAGGAGACATCTCAAATGCCTTCTTTAGACTTTGGT TCAAATAGTACCTAAGAATGAAATTACAGAAATTGAAGTAAATGCAAACAGAGTCCCAACA GTTGCAAGAAAATAAGAGACCTGAACAGACCGTGATGTGAAGGATCTTGTTGA
>Gm07_7 806619	ACTATATCACTGTCTTCTATTATATTTTTATGGATAAAAAATATAACATTATGATTATGATT CTTCATTTCTTGTTAATTGCGAGAGTCAGTAGCGAAATCCCCCAGCAAATGAAGGGTCG GTCTTTACACTTATTGCCACTTTCCATCATAAGCAGGGTGGTCTTTACACTTATT[A/C]CC ACTTTCAAGTTTCCATCATTAGCTTTGGACTTATCCTTCTTAATCCCACCTTTCTTCTCTTTCA CTTCCATGCGAAGAGATCTTGCTTGATTGCGAAAAGGAAGGAAGTGGAGACTTGTCTCTC AGAGATGGCGTTGCCTAAGGTTGATTTCGAGGTTGGAAGTATGAGGTTGCTGT
>Gm07_1 5490653	CTCTCCCCTCAACCGCTTTATAAATTGGGGTGGTGGCTTCGCCCTTGAAACTCGTTCTAGTG TATGTGATTGTTGTGACTCGTCTTCTTCGTCGTTATCTTCTTCTTTGTTGTTTGTGTGTTT GTTTTTCTCTCACCTGAAAATGTCTTGCTGCGGTGGTAAGTGTGGTTCGGAAG[T/C]GC CTGCAAGTGCGGCAACGGCTGCGGAGGTAAGATTTTTGTTGTCACTGTTGCTTCACA CTTCGAAAAATTGAGTTTTTCCGGGAAAAATTTCTTCTTGGGTTTTTATGGGCTTTTTTG GTACCCTTTTGGTTGAATTGCATGAAAAAATTATGCTTTGTTATTCTTGCATGAG
>Gm08_1 41320	ATTTCTCAAAATTTTCAATCTTATCCCACAATAGAGCCAACTTGTTAGTGTCTCTCTCGCTT CTAACACAATTCAACAGTCCCTTGCTTCTCCACTCTGGCCCAAAGCATGAATCTTGGG TCAAAGTAGCAAATTTGCTCCTCTCCTTGTTTTCTTCTCTACCAGTAGTAACCGTT[A/C]TAT CCTTCCACCACCTATCAAAAATCTCATATCTTCTCTCTCTACCATGCTGCATGTAATGGCC CTTGTTCTATGCATCCCCTTCCCATAATATTCAGCAATGTCTAGTGGCTCAACCAATAGC TTATAGAAGTGTGAAGTATTCACCCACTTGGCTCTCTTGTCAAAATCATGAGGTA
>Gm08_5 88567	AGCCTTTAGCTCAGAGTCTTCTACATTGCTATTAATGTTTCTAACAAAAAGTGTCTAGAAG GCTGTTCAACTAAAGGAAGTTTTCTTTAGAAGTTCCAAAGAAACCAAAATCTCCATCAAG AGCACTATTCTTTTTCTGAACCTTAAATGTTTCATCTCCTTCCATGCCCT[C/T]ACC ACTGCTAAACAGATCAAAATCTTCAAAATCATCATTTGTTCTGGCATGAGAAGTGCATCCT AACTCATCAACAACACCAGAAAAACAGATCATCTTCATCGGGAAGGAGATTGCCTATAGTGT CAGCCTCCATTTCTTCAAGAGATTTATATGGTTCTTCTTGGGAGGAGAACCAGC
>Gm08_3 461787	CAAAAACAACGACTGCACATTGTGCCCTCACGAGATCACCATCTGAGGTGGATCCTCAGC AGCTGAAAGATCTGTCAATTACAACCTTAGATACATATTCAACACTCCACACCCATTTTTGTA TTGCTCAAAGGAATAATTTATGTATATCAAGGAAGAGGAGTTTTAATTAATTCAATG[A/G]AT ATTTTTTAAGGAGGGACGCTGGTTATGATTGTAACAGGAGGTAAACATACATATGCTAATC ACATTTTTGAGCCAAACAAATTTATAGAAATATCTTAATTACAATTGGATGTAGTTACAATTT TCTACTGCTATTGTTTTCCGAAGTAGTGAACAGAGGATTGTGTTGCTGCATAT
>Gm08_7 669415	ATTTGAAGTCCACATATCCTGAAGTGCAGCGGCAGCCATAGCTTGGTACATATCTGATTGA TAATTCACCATTTGAGGGATCAAGCCTTGGCTGCATCCAAGGACTAACCAATACCCCTGA AAATTAAGAGATGGAAGACCTCTGTCTGTGTCTCGAAGCCACATTAGTGAAGAATTTA[C/T] TCCAAAATCATCATCCTTCAAGCCTGCTATGATCCATAGGGGAGAGTCAATGAGAATGAAT TTCCATTTATGCATAAATGATATTGACTTAGAAACATAAAACAGCTTTCTAAGTTGAAAATA GCTCAGGTCACCAGAAAAGTTCGTGAATATTGTCAGCTCAAATGTGGCAAGAAGCATA
>Gm08_1 3546289	TTACAGGGAAGTTGTGCTCCAGGACTACATGAACAACCTCTTTGGGAATGCTCATGCTGG AAAGAAGTCTCTTGATTTTGCAAGGATCAACTAGTTTGTATCAATAATATAAGAATTGAGAT GTTTGTATTTTCTTTTTCATTTTCATGTTTAATGAATAAAACCTTCAATATTGAGGCT[A/T]TAT GTCATAATAACTAATAAAGGTAAAGATATGGTTTGGTTAATCTTTATACACCTTAGCTTAA AGAAAGAAGAGAGATAAAAAAAATGAAAAATGAAAAATGATGGATGCGATTCAAAATGAAGA AAGAAAGAAGATGGAACACATATAGAATGAGTGCAGAATTAAGGTGTGACA
>Gm08_4 4999796	TCGATCTCCTCCAGGCGGTGGGACTCCGGTTCGACCAGGCCCGTGTATCAGCGCCTT CAGCAGGGTCTGACCGGCCAGGCCACCTCGTGCTTCTTACCCGCGAAGTCGATCG CCGACAGCCGCACGATGCGGTCCGCGACCTTCTCGTGGAGGAGGAAGAGGTGGAGGT GCG[A/T]GAGATCAGGGATTTGGAGAAGAACGAGACGAGGGGAGGCGGTGCACTTGCG AGGACAGTCTCTGCAGATTGCCACTGCCATTGGTGAATGTTGTTGCGCCAAGGAAGCCC GTCCAGAGTGCTCTTTTGGTTTCGGAATCGGAACAAAGGGAATGAATTTGGGTTGGGCC CGATCGTT
>Gm08_4 5844445	GACCGTTCCATTTGAACTGTTTGACGAGATGGAGCGGATCAGGTTGATCCGACCCAATCA TTATGTGCGCCAAACAAAAAATATATTATATCTTTAATTTGCACTTTTACCAAATTGACGAC TTGTGTAATACCTAGTCTCAGTGCCCAAAACAAAACCCCTTGCGTCTCGCCGCACA[A/G] TTACACTCCGCAGCACACTGCAATCGCCACCACGGACCAAGCTAGTTCGCCGCTCGACG

	TCGTCGGCCACCGTCACGTCTCGCCGTCAGGTGCCTCCCTCTGTTTCTTTGGGGTTCTGG GCATCTTTGTTTATTTGCCACGAATCTCTCTTTACCAACTCTTAACTAAAACTTGCTGATT
>Gm09_3 799409	CGACCATAATTGTAGTTGCATCAGCCACATTTGTCAGCAATTTCCCACAATTGTGATCATA AAACCTCGACCATATCAAGAAGAAGAGAGCTTGATAGCTTTGAAAGCAGGATGAAGCAAT CCAAGCTCATCCTTAACCTCAAGAGGATTGCTAGTATCATTCCCTCACCCTCTTCACCT[A/C] ACGGCTACCAAGTGACTTATGAGTGAATCCATACATCTGCAAAATCTGGTTATCCCCAAAG TTATAGGCCCTATCCATCTGCTTCAAGCACCTCTCAACAGGGTAATCCCCAAACTTCCCAC AAGGCTTGACCCCCACAAAGTGAGTCACCAAAGGCCACCTATGATCACCAAGCCCCGG
>Gm09_3 799416	AATTGTAGTTGCATCAGCCACATTTGTCAGCAATTTCCCACAATTGTGATCATAAAACCTC GACCATATCAAGAAGAAGAGAGCTTGATAGCTTTGAAAGCAGGATGAAGCAATCCAAGCT CATCCTTAACCTCAAGAGGATTGCTAGTATCATTCCCTCACCCTCTTCACCTTACGGCT[T/C] CCAAGTGACTTATGAGTGAATCCATACATCTGCAAAATCTGGTTATCCCCAAAGTTATAGG CCCTATCCATCTGCTTCAAGCACCTCTCAACAGGGTAATCCCCAAACTTCCCACAAGGCTT GCACCCCACAAAGTGAGTCACCAAAGGCCACCTATGATCACCAAGCCCCGGTGATAA
>Gm09_6 112681	ATGCATTGAAGGAACAACAAAATAACTAGAAAGGTATTGAGTTCAGAATAACCTGTGAAAT AAACGTGCATATGGTTCTAGAGCCGGAATCCAGCAACGAGATGCTCATCCAAAGCTGTT GTTGGAGGAGGAAGTAGGAGTGCCGGGACAGCAGCTGCTGTTAAAGTAGCAGTCTCAT[G /A]ACGAGCAACTTCATCATCACAACACTTAATATAACACCAGCACCATTAGCAACTGCCC ATCTTGAGTTGATGGCAGGAGTTGAGGATGCTTCCAGAACC GCGAGCAGAACTAAAG AAAAACAAATTGAGTTAAGAAAAGATTATTATAAATCTTGCTTTTCGAGAATCAAAATGAT
>Gm09_1 5218066	CATCAAATCCATCTGTATTTGCGTGTAAGTGAAGGAGCTGAGCTAGGATGCCGGTCCTTGT AAGGGTATGATCTCTCTGCTTTAAGCATCATGCGATTTCTTTCAGGTTCTTGCTTTGC ATCATGCTCTCTTTTATAAAATCCCGGCTCAATTCATCAGGGTGCTGTCTGATGCTA[C/T] CCAGATGTGCTGATTGTGGATCCTGAACACTCTTCTCTCAAAACCCTCTTGCCGCTTCTG GTTATCTCTACTGCTTGCTGATCTTGCTTTGCTGTTATCACTACTTTTTGCAGCCTTTGAGT CCTCCCAATCATCATCAACTTCATCAGGCAAGGTGGCTCAACTTGAGATGTTACTT
>Gm09_1 7990620	AACATATGAAATTGATGGAATTAAGGGTCAATTAAGCCCCATAATACATCAATTTTTATTTT TTAGATAGCATTAGTAGAGTAAACATGAGCTTAATTTGATGATTTAGTTGTGCCCTGAG TAAGGTAAGCATCAATTTTTTTGCTAACCAACAGCCAGTTGCAACAAGGTATGA[A/T]CG TCAGGAATGTGGTCCTCTTGTTTCTCGTATTCCATTGCCAAAGCACTACGCTGCCCTTTC CTTTTGAGTAACCTGCAATTAGACTTGAAGCTCTTGATAGAGTCCAGCACGTCACCCTC TATGACCTTCATTGTCATTTTGTTATTTTCTTGCTATGCCTTCAAGCACTTGCT
>Gm09_3 8634550	AGGTGGTGGCGGAGGAGGTGACTCTATACAAGGTGGTGGAGATGGGGGTGGCGGTGGT GGTGAATAAACTGGTGGGGGAGGAGAGTGAAGTGGAGGAGGTGGTGGCGGTGATAAATA GGGATAAACTGGTGGTGGTGGTGGTGGTGAATGTGGTGGTGGCGGTGGCGAATGAGGT GGAG[A/T]CGGTGGAGGGGGGAATACACTGGTGGTGGAGGTGGTGGCGAATACACCGG CGGTGGGGGTGGTGGAGGGGAATGAGGAATGTAACAGGCGACGGCGGAGGCGCGAT GGCAATGGTGGTGGAGGAGGAGGAGGAGGCAATGAAGGCACAAAAGGCTTACACC TAAACGAATTACA
>Gm09_4 4037531	AAAAGTACACCAAAATTGAACAAGGATGAGGATGATGATTTGTGGGGATCTATAGCAGCT CCTGCGCCAAAACTGCAAGACCTTTAAATTTGAAATCAGCTCAAAGTATGATGATGATC CTTGGGCGGCCATTGCTGCTCCAGCACCCACGATCAAGGCCAAACCCTTATCGGCTGG[A /G]AGAGGTAGGGGAGCCAAACCTGCTGCTCCAAAATTAGGTGCTCAGCGGATAAACCGG ACATCATCAGGCATGTAATCCAAATTGCCTGGGAACCGAAGGAATGGCGAAAGAGTTAAT CTTAGTGCTAGTCGCCCTTTCTTACGGTGGCAGTGACCATGTTAAATACAAGACCCGTT AT
>Gm09_4 4175240	GACAGGAAGGTCAATAACAAGTCCAATATCCAATTGGTAATCAATGTCCAATCTGCACCA ATGTCCTTTGCCAAAGCTCAGTAATATGCTATGAGGCACCTTCACTGGCACATCCAACATTG TTGTGTCACTTGCTTCAATGACCCTGGGTCAGGTATTGTTCTGCTGCTATCTCCC[G/A]T CACCCTCAACACAAAAATAAATAAATAATTAATAAATAAATAAATAAATAAATAAATAAATA TAAAGTAACAAATAATCAAGGACACAAAAATAATGTAACCAAAAAATTAACGATCACACG AAAAATGCAGATTTTTCAATGGATATAAGGGAGTGGATTCTTCCTTTATGCAAT
>Gm10_4 962619	GGAGAGAAAGGGAACGAGTTGTTGAAGAAGAAGGTCTTGAAGGAGAACGGGACGGAG CGAAAGAGAATCTGAATGTGGAGAAGCATATTGCAGCAGCAGAGGTTTCCGCCATTGGAA CGGTACGGAGTTAACACAAAACGTTACGTTACGTTATCAAAACCGCTTTCGATTATTCAT[ A/C]CAACTACTCACCGCGCTTCGTTCCGACACCCTCCTCCCTCTTTGTCGCGCATTTCT GGATTATTTTTTTATTTATTTATTTATTTGCTTAGAGTGATGTTTTTTTTTTTTTTCT

	GGCTGAAAAC TTCACATGGATACCGCACACACGTTCTATTTTTTATTATTATTATTA
>Gm10_3 3420456	TAGGTGGCGCGGAGCAGGCGGAGGATCCGGGGAGAGTCGCCGTGGGAGCCGGAGATG AGGTAGGCGAGGGAGGGAGGGGAGGGGGTGGAGGGTTTGTGGGTGTGGAAGATGGGG TGGAGGAAGGGGTCCGGGCCGAGTGGGGAGAAGGGGAGGTGAAGGAGAGGGAGAGG ATGAGGGT[T/C]AGGGAGAGGAGGGAGAGGAAGAGGAAGGAGTAGAGGAGGAGGGTGG GTTTGGGCTTTGGATCTCTGAGGTTGAGGAAGGTGGAGAGTGAGGGCCAGTGCAGCTGG GACTGGGAGGGTGTGGTGCTGTGCTGTTCTGCATTTTTGGGTGTGAAGGATGTGATGAT GGATGGTGGTTTGCT
>Gm10_4 7189583	AGACAAAAACAACATGCAATGAGATTGTTGGCCTCACCGCCGACAATTACATAGTATATAT TTCATAAACATTCAACCCAAAAATCATTATAGAGCATGGCCTTTACTTAATTATGATTATTA GTAGTAGCAACATAACATAACTCAAATGGTTGGTGATTTATTTTTGGTCCAGTAT[G/A]TG AATACAATGCCAAATTCATAATGTGATGCGAGTGACACATGCCATTTACAAGACATGCAA CGCATCATCCCCTATTGCTACCTTCACCACTGGCAAAGACTCCATCAACATAACCAATCAT GGCCACCACCTTCTTCTTCTGCGGCGTCCCTGGTCACTGCCAAGCAGGACAGAAGGT
>Gm11_5 650369	CCTTCTTCGCGGCTTCGGCTAACTTGACGAAGCCTTGATCTTGACGAGTTTCCCTCTCG CAGCTTGGTTCTTCAGTTGCAGACCTAGTATCTTCTTGAAATTCGCAGGAAGCACCGCCTT GTGCTTCTCCTCCATGTACTTCGCTATCGCATATGGACTCGATCCTCTTCTCGTT[C/A/C] GAGCAATCAAAGCGTCCTTAATCATCTACAAACAATAAGACAACCAAAAAAAAAAATCAACA AACAAAAAATAACAATCACGTAATAATTGTGGATTTCATGAAAGAAATTTACGCGAAATTAC CTGGAGATATGGAGGATGAGAAGCAGTTTTGGCCTGTTTCGGCTTCTTTTCTTCTCGG
>Gm11_9 563735	CAGATAGAATTGGTGAGCAGATCTCTTCTCTTTTTAAAGGTTTTTTGTGATGCAATTA GATTAGTATTTAGTCACTCAAGTTGCTCAAGCTAATAAGCTCTGGAGCAATATGCTGAAAT TCTAGTGTGTGCTGTGTTACTTATCTCATTTTTCTTTTTGAATTGTTAGGTTCCAT[T/C]CCT TAGGGACAATTATGCATATCTTTTACACGATGTGGATACGGGGACAGTTGGTGTTGTTGAT CCTTCTGAAGCTGCACCTATCATAGATGCGTTGAGCAAGAAAGATTTAAATTTAACTTACA TAATGAACACCAACCACCATCCTGATCATACTGGTGAAATGCAGAGTTGAAAGA
>Gm12_3 433264	ACGATTTCCCAAATCATAATGATTTATGAATACACGACAGCTAGAAAGACACCACATCAA TCCTCTTCATCTTTTGTCTTCTACATTTCTATGAAACCAGAGGACTTAAACTGAGATTTTATT CCACTTTTTAACACATGCTAACTTTAGGTCTTAGGACAAACATGAATCACAAAAA[A/C]TG GCCGGGTGTTTATAGATATCAAGTTGCAACAAATGAATGATACAGAAGACACTTGGTTTAA TTCTCACATATCAACCAAGTCCAAATGGTGGTGCCTAAAAAAGTATGACCACCAGCGCCC AGATGGGATAAACCAACAGAAGTATTATGCCAATGGAATTTGATATGCCGTGAGCTT
>Gm12_3 862466	GGTGGTGTAATCAAGAAGGAATCAATTATTACAACAACCTCATCAATGAGCTATTGGCTA ATGGTCACAACCTCACTATTCAAACCTCTGTTTTTAAATTACATTTAAATTTTAGCATTTAAT TATTAAGTGTTGTCAACAAAAATTAATCAAATATTGTTTGTCAATTGAGACAATTA[T/C]AGGT GACTGCCATATGTGACTCTTTTTCATTGGGATCTTCCCCAAGCTTTGGAAGATGAGTATG GTGGTTTCTTAAGCTCCCATATAGTGTGAGTCTCCATTTTATAACTATCCAGCAAAATTAAT GTTTTCTCATCATCAGTAGATTGTAGTTTGTAGTTTTTACTAAAAATGTTT
>Gm12_1 0340760	AGAGGGAACAGGATGCATAGATTGTAATTGAGTGTTTGTCTTTCTTCTTACAGGGAAGAG ACCGAGTCAGGTGAGTGTGAGACAAGGGATTCTGTACAGCGATTCTACAGTGAAGAGAG TGAGAGTGACAACTATGGAGGTGGGATGGAACCTCGTCAGAAGAAGGAGGATCTGAGC[ G/A]CGATTGTTTGTGGCACATGAATGACCGATTGGGTCACCTTTACTTCCAATATTTTGA AGATCAACTCCATATGGGAGAGTTCCTCTAATGGATAAGGTGCGTTTGTCTTCTTGTGTT GGCCATTTCTAACTTTTTTTTTTGCAGCTTTATTAGTTATTAGATGTGCTGAAAGAAATTC
>Gm12_1 5607093	TCTCCCTATGCCATAAGCTTTGTTCTTTTCTGTAAACATGATTCTCATACATCCACGTACA GGTGTCAATGAAAACCTGAGGTAACCTTGCTTCGTGATGTGCTTCGCGGTGATGATGTGTT TGAGTTGCAGGTAACCTTCTCAAGCATGTCCAGAAGAGTATCCATACAAGGATGA[C/T] GAGTGAATTTGTTTTGTAGTATTGTATACTCCAGGTTTTTCACTTTCATCATCTCTGCAT TATTAGATTATCATATTCTGACTGGTCTGTGTATTATGATAAAACACTAAATACAACAG AAAATCTTTAATTAATGATTATCATATTGGCTTTGACAGCGAGCTACGATCTATT
>Gm12_2 4557740	ATTTGGGGCAGTCTCAGGTGTAACAGCCACCGGATACTTAATGCTTGAAGGATTCACCGA CACATTTAGGACCGAGCTGTTGACATGAAGGTATTTCTATCATTACCCCAAGTCCTACCC AGGGTGTCAATTTGGGCAGTAATCAAAGGACCCCCATGTTCAAGCGATAAACAGTTT[A/T] ]AAAGGCAAGTTCAGAAAGTCCACTGAATGTTGCTGGCGGGTTAACAGCCAGTGCATGAT CAAAAAACAACCTCATTCGGCATTGACACGACTTCAATTGCATTGACAAAGGCCACTGAACC ATTCAAGGAATGAAGGTGACAACCAATGTATCAGAGGTAACATTGATTGCATACTCCTT
>Gm12_3	CATATGTGTATGAAAGAGTTTATGTCATCTGACACCGCATGTATCAGTTAACTTTTACTAT



5462187	GAATCAAAGTGGTACCTGAGAGAGCGAGCTTGGTCAAGAGACCAATCAATTTGAGTAGTT GGACCTCCTGGGGGAGGACTAGAAACACTTGTAGGTGAGTTACTGTAATGAAGAATG[A/T] AGGTAGCAGTCAACACTTGGGAGGTGAAACGTGTTGAGACCACAATATAATAGTCTTGAG GTGGTTGATCAGCAGTTACTAACACCGAGTAACTCTGCCCCAAATGAATGTCAAGGGAAT CATAAGTGTTTTGGAGAGTATGGATTCTTCCACCTCAACTAGCAGCATCTTATGCCCTT
>Gm12_3 6713901	TCTTAGATGGTTCACTCGTCAAATTCCTTGCTAGAGCATAATCTTTCCTTTTTGATCTTT GTAGCAATTCCTTTCTCTTGATTGGCTCTTTCTTTCTTCTCTTTTATCTTTATTCGGATT TGCCTATTGTTGTGATTCAATTGGTTTAATTACGATCTTCAACCGTTTATATTTG[A/G]TTCTT TCTTGATCTAAATTATACCTCTCTCTCTCTCTCTCATTCTGTCTTAAAAAAAAAAATTTGA TGTTGGTATTTCACTCATGTATGACAATCTACATGATATGATGAATAGCAGCTCAGAAACA ACCGTTGACTCACTTTTAATTTTTCTCCCATGAGTATCCATTCTTCAAC
>Gm13_3 5617464	GTGATAATTGCTGCACGGAGCAACAACGAGATGGATGCATCCAACTGAAGAACGAGTTC CCTGAGCTTCTTTCCATCAAGGAATCACTGATCAAGTATGTCTTTGAGCCAAACAAGAAAA CTGCTTAAATATATTAAAGCGGTTTCATGTTTTAGACATCACAGATTATGCATTTTTG[A/C]G GCGGGCTACATAGATCACTCATTATTTCAATTGTTATTTTGTGGGTTTTGATAAGAACCATA CTACAAAAGCTAGCCATGTAGTTCCTGTTCTCTTAGCTATAAGCGATTCTGTTTCAGCAT ATTTCCATTATATTACCGTGTTTAGAGGGTGACAAGTATTATAACTGGGCCAGATA
>Gm13_3 5823484	AGCCTCGTCCCCGACGTCATCGATCTCAGACCTCAAAAAAACTCAAAAGGGTCCTCCTCA AACCTCCCAAAGAACACAACCTTTGGACCAGATTGATCCAAACAAAGTTAAATAGCCTTCTG GGTCGGTCACAAATTGAAAAGGGTGTGTGTGTGTTAGAAGAAGAAGGTTGTGTTATAA[T/C] JAGAATGTGGAGGAGATGGTTGGAAGTGGCAAAGTCAGCACATTTGGTGGATTTGGGAA GTGGGTTTGTAGCCGGAGAGTGATCGGAGATCGGAGGAGATCTGGGTCGGAGAAATGG GAAATGGTTTTGATGAGAGTGTGTGTGGATGTGTGTGAGAAAGAAGCGGAGGAGAAGAA GAAG
>Gm13_3 5823512	GACCTCAAAAAAACTCAAAAGGGTCCTCCTCAAACCTCCCAAAGAACACAACCTTTGGACC AGATTCGATCCAAACAAAGTTAAATAGCCTTCTGGGTGCGTCACAAATTGAAAAGGGTGTG TGTGTGTTAGAAGAAGAAGGTTGTGTTATAAGAAGAATGTGGAGGAGATGTTTGGAAAGT[G /A]GCAAAGTCAGCACATTTGGTGGATTTGGGAAGTGGGTTTGTAGCCGGAGAGTGATCGG AGATCGGAGGAGATCTGGGTGCGGAGAAATGGGAAATGTTTTGATGAGAGTGTGTGTGG ATGTGTGTGAGAAAGAAGCGGAGGAGAAGAAGAAGAAAAGGAAAAGAGTGAGGAGGAGG AGGC
>Gm13_3 5823533	GGTCCTCCTCAAACCTCCCAAAGAACACAACCTTTGGACCAGATTGATCCAAACAAAGTTAA ATAGCCTTCTGGGTGCGTCACAAATTGAAAAGGGTGTGTGTGTGTTAGAAGAAGAAGGTT GTGTTATAAGAAGAATGTGGAGGAGATGGTTGGAAGTGGCAAAGTCAGCACATTTGGT[C/ T]GATTTGGGAAGTGGGTTTGTAGCCGGAGAGTGATCGGAGATCGGAGGAGATCTGGGTC GGAGAAATGGGAAATGGTTTTGATGAGAGTGTGTGTGGATGTGTGTGAGAAAGAAGCGG AGGAGAAGAAGAAGAAAAGGAAAAGAGTGAGGAGGAGGAGGCCTTGATTAGCCAAATT TGCG
>Gm13_3 5835159	TAGTTGCCGTACATTGCGAGGCCGGAGAAGATCTTCGCCGCGATGGCTCTCCTCCTCCTC TCTCGCCTCTTGTTGTTCTCCCGCTCCTTCCACGTTGGCTGCCTCGCGACGGACGTCATC CTCTTCTCGCCGGA AAAAATCAAAGAAGCTCCGAAAAGAACAAAGGAACAAAAAAGAAA[A /T]ACACACACGCGAACACGAATGTTAGAAAGAAACACAGCACACAGCGCGAGTAAGCGAA TTATAGAGGAAACGACATCCTTTCTGCTTTTTTTATATAAGGTGAGCTAAAACGACGACGT CGGATTTGAGAGATTTGGGATTGTGATCTGAGTAGATCTAGGCGAGGGAGGTTGGGTGG TG
>Gm13_3 5862124	ACTAATTAATGTCTTATGCTACCAAAGCACTCATACAATCACTAACAATGACGAACACTCA AAATCACTAGTCTTAATTA AAAACATCTATCAAATATGTTTTCTGAACTTTGCAATTTCTAAT TCTCTAATTGCCCAATTTCTTTGCCGCGTTAGCTTCAACCCTAGCCTCGTCATTT[A/T]TTTT GCTTCCGTTGTTACCATTGCGCATTTTTGAACCACCTCATTATGTTGTCCACCTTTCCCT GGCTGATGGGTGACCCTCTTTCCCTCCAACGCTTCCATGGCTGGGGCCTGACCCTCTTG TTCTCCGTGCGTTCTACTCTAAACAACAATAACGTCAGTTCTTAATTCATAT
>Gm13_3 5862205	AACATCTATCAAATATGTTTTCTGAACTTTGCAATTTCTAATTCTCTAATTGCCCAATTTCT TTGCCGCGTTAGCTTCAACCCTAGCCTCGTCATTTCTTTTGCTTCCGTTGTTACCATTGCG CATTTTTTGAACCACCTCATTATGTTGTCCACCTTTCCCTGGCTGATGGGTGAC[A/G]CTC TTTTCCCTCCAACGCTTCCATGGCTGGGGCCTGACCCTCTTGTCTCCGTGCGTTCTCA CTCTAAACAACAATAACGTCAGTTCTTAATTCATATTTTCAATTTTATATATATGACAGAACAT CTAGCTAAGGCTGAATGAGATTTATGCATGTAAGAGTGTTTAACTTGATCACT

>Gm13_3 6473402	CAACAGGAAAGAAAGCCCTACAATAGAACTATTCATTAACCCAATGAACAAGCAATGTGCA TATACAATTACTGCTATGATTTAACTCAATGTGCAAAGTTAGTTTGACTCTGCAAGGCTCA ATAAACAGTCCTAAAACCTAACACCTACGCTTTATAGACATCTCCGAATGATCCCT[G/A]TC CAATAAGCTCCAATGAAGTGAATCTCGATCCTGAGGCCTCACTAGACCTGCCACATCAG CCATTCTCTCAGCAGCAAGTACTCAAAGCCTCATACACAGACTAAACCACATGCACCTGA AAAACCAAAGAGCACAACCTCAGAGATGTAAATAGTGAAAATCACAATCTCAAATCTC
>Gm13_3 7946607	GGGGAAAGAGATACTCCATTTGTGCTATATTTATGGTATGTTTTATTTGTGTGGTGTCTG GAAATGGCCTATTTTCATTCCCAATCCGTTTCTCCAGTCTCTGGGGGATTTGATTGATGAG AATTGCTTATGGTGGGTGTATTTCTTTTCCAATGAAATGCTACAACATTTACTGCA[T/C]T GTTGGAGATTGATTCTTCGATTGTTTAATCTTGACTTCCATTCTGACCAATTCTATTGCCGC AAAACATGGAACAAGTAAATAAACGGGCTGAGAGAAAATAAGCATTGTTTCACCAGTCAG GAAAATAGACTTAACTTTCCTCAAAAAATAAAAAATGGAAAATAGACTTTACTCCAT
>Gm13_3 7946608	GGGAAAGAGATACTCCATTTGTGCTATATTTATGGTATGTTTTATTTGTGTGGTGTCTGG AAATGGCCTATTTTCATTCCCAATCCGTTTCTCCAGTCTCTGGGGGATTTGATTGATGAGA ATTGCTTATGGTGGGTGTATTTCTTTTCCAATGAAATGCTACAACATTTACTGCA[T/C]G TTGGAGATTGATTCTTCGATTGTTTAATCTTGACTTCCATTCTGACCAATTCTATTGCCGCA AAACATGGAACAAGTAAATAAACGGGCTGAGAGAAAATAAGCATTGTTTCACCAGTCAG GAAAATAGACTTAACTTTCCTCAAAAAATAAAAAATGGAAAATAGACTTTACTCCATT
>Gm13_3 9111783	TCCGAGATGTACTCCTTAAGGTTAATATACCAATTCTAGAGATCTTTGGCTTAACTCTTTG GCTTCTGTGTACTTAAAAGTAGATGATTGAATAATTAATTCAATATAAAAAAATATTGATTTG ATTTTACATAAAAGTTAAAAGACATAAAATAAATAAATAAGAAAGGCACATGAT[A/G]GTGT GAGAGGAAAAATGCATTAAATTGAGAATAACAATGAAAAAGTACATCAATACAAAAA AGACTAAACATACATTAACCTGCTGTTGGTAGCACAAGAAATCTGCTGCTTATTGCAAG AGCATGATCTACCAGCAAAGTTGAAAATAAGAATAACAATGGAACCTACTTATT
>Gm13_4 0270504	TTTGGTGGCACATTCAATAACAGCTGACAGTTTCTTCCAACCTGATTTATAGTATATCTCAAG TAAACTCCTTGACAGACTTAGGTTGTTCTGAAGCGTGCCAAAACCAACCGGTCTAATAGA GACATCACATGTAGCAGGTAACCAATCAGGACCAAAATTGGTCTCCCTCGCTATTGTA[G/A] CTGATAGACGACCAAAAGATTGGTGAATAATACATTTTTGACATGCCATTTTGAGAGCATT TCTAGTCATGTACATACATGATCCTCCAGACTTTCAAGCCTTTCCAAGGATACATGCAAAA AAGAACTAATGACTGAAATTTACATTCTTCAACTTCATTAAAACCAATGAATCAAAG
>Gm13_4 0270505	TTGGTGGCACATTCAATAACAGCTGACAGTTTCTTCCAACCTGATTTATAGTATATCTCAAGT AAACTCCTTGACAGACTTAGGTTGTTCTGAAGCGTGCCAAAACCAACCGGTCTAATAGAG ACATCACATGTAGCAGGTAACCAATCAGGACCAAAATTGGTCTCCCTCGCTATTGTAT[T/C]T GATAGACGACCAAAAGATTGGTGAATAATACATTTTTGACATGCCATTTTGAGAGCATTT TAGTCATGTACATACATGATCCTCCAGACTTTCAAGCCTTTCCAAGGATACATGCAAAAA GAACATGACTGAAATTTACATTCTTCAACTTCATTAAAACCAATGAATCAAAGT
>Gm13_4 1274238	TGGATCCACTGCATGGTTGTTTGAGGATATTGAGGAATCAACGCTAGAGGGTGTTGGTGA TGACGATGCAGCTTCTCCTCGCTTTTTCTCCTGAGCCTTTTTCTCATCAGAGTCCTCCAG TAATTCCTTGATCTCATTGTCAGTGCGCCCTGGTAACCTTGCGAGCAATTCCTGACCACC[A/T] GCATGTCTCAAAGTAACCATGATTAGATTTGAAATTAACCTGGATAAAGAATATGAAACGCA AACTTAGATGTAGTCAAGTGTTGTTTTCTGATCAGAATTTGGAGTTTTACCATGGTGATCT GCATAATAAAGGTTTGCAATTAAGATCCATCGACATAAGTCATATATAAAGGTGAAAC
>Gm13_4 1274239	GGATCCACTGCATGGTTGTTTGAGGATATTGAGGAATCAACGCTAGAGGGTGTTGGTGAT GACGATGCAGCTTCTCCTCGCTTTTTCTCCTGAGCCTTTTTCTCATCAGAGTCCTCCAGT AATTCCTTGATCTCATTGTCAGTGCGCCCTGGTAACCTTGCGAGCAATTCCTGACCACCT[C/T] CATGTCTCAAAGTAACCATGATTAGATTTGAAATTAACCTGGATAAAGAATATGAAACGCA ACTTAGATGTAGTCAAGTGTTGTTTTCTGATCAGAATTTGGAGTTTTACCATGGTGATCTG CATAATAAAGGTTTGCAATTAAGATCCATCGACATAAGTCATATATAAAGGTGAAACT
>Gm14_1 069974	CACCTTCTTTGCATAGGGCAACCTCCTTTCTCTGTGACTCTGAGTATCGTATCGTTTTATTA TTATTTCTAATTCTCTTTATCAGTATGTATTCAAACCTAGCTAGGTTTTTGAGCTCCC TTTCTTTCTTTCCGATCCCACCGTCAAACCTTGTCATTCTATATGCTTTGCAGGGT[A/C]CATT GAATATGTCTCCGTCGTTGGATCTTCTCCCAAGGGTGGCTTCAGTTTTGATCTATGCCGA AGAAACGCAATGCTTCAAACAAGGGCCTCAAAGCCCAACCTTTCTCAAACTGGCACC ACTATTGTCGGTTTAGTTTTCCAGGTCTGAAATCATTCCCTCGCTCTTTTCATGT
>Gm14_4 838692	TTTTATGCACTAGTACCTGGTGCCGTTATGGTGTGACTCAGTCAACCTCTTAAATGCGGGT TCCGAACAAGGAATTGGCAGTTTTCAATGCACAAGATACTATTACTAACCAGCAAGCAAG AAAATGGAGAAACGGCTATTAATAACAAAGGAAAAAAAAAAAAAAAAAGGTTATGTGT[T/C]T

	GGGGTGGTAACAAGTGACCAAGAATGAGTCCTTTGGTGTAGAGATCTGACAAAAGGAAGT ACACCTGAGAGAGTAACTTGGAGGTTGGAACATGCTTGGGACTTGATTAAAAGAAACA ATATAAAAATGAAAAGAAAGCAATGGTAAAAAGGAAGAACTAAGAAGAAGGGTGTTTAA
>Gm15_5 573351	GAGGAAACCTTGTGTTTAAGGATGGGAAGCATGCTCCTGCACCGTGTGGTGTTCAAATAC TAGCCAAATGAGTTAGTTAAATACTGGTGATCCCCATAACTTTCCAAGCTCATCTCAATTTTC ATTTGGTTCTCTATGAATTCATGATATGCTTATCATGATCCTTATTCATGTTGTAGA[A/C]AG GTTTCTTTCAGATGCTAGTTACGTTGGGACCTAAATAAGAGATGGAAGTAGATGCTGGCAT CGACCACTGACCTAGCATTCTCCTTCTCATAGACTAGTGCATTACCAAATGTATTATTATC GTTTGAATGTTTGGACAATGACATAATGTTGTTCTTTTCTTGAATGCGTATGTAGG
>Gm15_7 814112	ATAGGGCAGAGAAAAGCGAATAATTGAAAAAATTAAGAAGAGATTGGGAACCTACTCAGTT CCTTCTGGCGAAGGATCCGAAACTGCTCGGGGGAGAGAATGACCTTCCATTCTCTTCA GTTTTGTGAATCGGCGTTGGTGCTGCCATTTTTTCCAGATGCTCCACACTGAAAATAA[T/ C]AAAAAAAAAATCCCTTTTTACAGTGCCCAATTGCAAGGAAAAACAGGGGAATCTTCCA AGAAGATTTTCCGGGAAAATGAACCTTGACCGGAAAATTGGGTCTGGACGCTGACAGTGG GGGCACGTGGCACGCTCATGAACATAAACTAAGAAAACGAAAAACGATTTTTGGTTAAGA AA
>Gm15_9 432381	CAGATCCGAAGGAGGAGAGAGAGACCCAAATGGAAAAAGAGTTAACTTGAAGGGTTATG GAGAAAACCCAGGTTGGGAAGGGAAGAATTGGAAGAGGGGAAAGTAGGAAGAGAACAAA AGAGTACCTAATTTGAAGGATCAGAACACAGCTTCGTAGCTTTTGCAGAGAGAGAGAGAG[ G/A]AAGAGAGGGAAACACTGTAGATAGGGAGCTCCTGGTAGCTTAGGCCTTAGAGAGAG AGAGAGAGAGGGGTCAAAGGGTAAAGAAAAGGCTTTTTTCTTCATTCAATAATTTCAATGGT TTTTGTGCGCCAAAACCAAGAGAGGTCCAAATCACTAAACAGTTCTGAGGTCTTGCACCTG TC
>Gm15_9 432383	GATCCGAAGGAGGAGAGAGAGAGACCCAAATGGAAAAAGAGTTAACTTGAAGGGTTATGG AGAAAACCCAGGTTGGGAAGGGAAGAATTGGAAGAGGGGAAAGTAGGAAGAGAACAAAA GAGTACCTAATTTGAAGGATCAGAACACAGCTTCGTAGCTTTTGCAGAGAGAGAGAGAGA A[C/T]GAGAGGGAAACACTGTAGATAGGGAGCTCCTGGTAGCTTAGGCCTTAGAGAGAGA GAGAGAGAGGGGTCAAAGGGTAAAGAAAAGGCTTTTTTCTTCATTCAATAATTTCAATGGT TTTTGTGCGCCAAAACCAAGAGAGGTCCAAATCACTAAACAGTTCTGAGGTCTTGCACCTGT CTG
>Gm15_1 5214016	TTGTTAATAATATAATGCCAAGAGGCATGAAGAATTATTTGTAAACAAAAACAATGTCTGAA ATTACCTGAACACAAAGAACAGCAGCTTTGTGACCCTCAAACCTGTGGATGGGTGATCCA ACTCCATTAGTGGTGAGATTGCGGCGATCAAACATGCGAACAGAATTGTCTGCTGAC[A/T] TGCAGAGATGTGGCGACAAGGCAAGTGGGGATGTCCAACTAGTGTACATGTCCAAACTA AAATGATGAATTAGTGACAATGAGAATATACAACAGCAGAACATACCCGGTAAGAATCAG ATTATCATCATGTGGATTCCAGTCCACAGCTGAAGATCAGCATTATGAGCTTTCTCAA
>Gm15_1 7582711	CATGGTCTGCATTCTGTCAAGCCAGCAACTTTTCTCTTTTTTGACAACCCGTTTTTCTTC TCAAATATCTTTCTCTTCTCGATTTGCTTTTGGAACTGACTGTCCAACCAAGGATGGGGGT GTTTCAATGCAAAGCTTTTTGCTACGTGCCCCAAGTGGAGTTTCTTGATCATAAAT[A/T]CT CTTTTCAGCTCACCGCGATGGGCTGTGTATGCACGAACCCAAGAGCAAATGCCTTCCTG GAAAGTTCATCCATCTCGGGCTGAAAAAGGACAAGGCATAAATAGGTTTATCTGCAAGGG ATAACTTTTAAGAAATCCAATGGTAATGATTCTACAGTTAACTTTGATGACATGTTT
>Gm15_4 6990101	NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNTGTCCGAAGACGACTTACCCTA CCATCTCCGACTCACCCCCAACGGCGACTTAACCACCGTCGGCCGTTACAACTTTAACGG CC[A/G]ATTGAAATCCACAATGATCGCTCACCCCAAAGTCGACCCCGTCACCAACGACCTC CACGCGCTCTCCTACGACGTCGTCCAGAAGCCTTACCTCAAATACTTCCGCTTCTCCCCA AACGGCGTCAAATCCCCGACGTCGAAATCCCTTTGAAAGAACCCACCATGATGCACGAT TTCGC
>Gm15_4 6990104	NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNTGTCCGAAGACGACTTACCCTACCA TCTCCGACTCACCCCCAACGGCGACTTAACCACCGTCGGCCGTTACAACTTTAACGGCCA AT[G/A]GAAATCCACAATGATCGCTCACCCCAAAGTCGACCCCGTCACCAACGACCTCCAC GCGCTCTCCTACGACGTCGTCCAGAAGCCTTACCTCAAATACTTCCGCTTCTCCCCAAAC GGCGTCAAATCCCCGACGTCGAAATCCCTTTGAAAGAACCCACCATGATGCACGATTTT GCAAT
>Gm16_2	CTTCTGACAAACCTTCAGTACCATGATTCCAAGGTTCTGGAGCACGCCTCTGTCTGTTTGA

5690145	CTCGAATAGCTGAAGCTTTTGGCTCAAATTAGATGAATTGTGCAACCATGGACTTGTAACA CAAGCTGCCTCCCTCATATCTAACAGCAGTTCTGGGGGTGGTCAGGCTTCTCTCAGC[A/C] TGCTGACATATACTGGTTTAATCCGACTTCTTTCAATCTAGTTAGTTTTATTTTGGATTTTCT TTTCTATATATATATTCATTTCTTGGGGATATGTATATATTTGTACTATGTTCTTGCAGGAG GAAATTCATCGGAAATCAGTCCTTCCATGGAGAAAGGAAGAAAAGGTTTTATT
>Gm16_3 0464934	TCTCTCCCTCCTCATGAGGAGAAAAAAGAAGCTTTCAACTTTCTTATTATTCATCCAACAC CATCTCAAAATCGAAAGGAACAAGCTTCACCATCATCATCACCTTTTCACTCCTCATGAG TACCAACCAAAACACCACCTTCTTCCCCCTCCACATGGCCAAATAACTACTTCTGA[T/C]T TTGTTAATAGCCAGAACAATCCATCCCAGGCTACAAATCAAGGTAAACTATATATTATTGC CAGGTTGATTATATATCATATTATTCTAATAATTCATATTACCATCGATGTTATTATATGGTC CTCTCAATTATATATATATCATATCATATCAATAATCTGGATCCACATCT
>Gm16_3 0539483	GGTTTGCAGAACTGCCAGACTGTGTTGTGCCAACCAACAGGCGGACGGGCAAGGTTAA CCGAAGGGTGCTCTTTTAGGAAGAAGGGAGATTGAATGGTTTGTTAACAGACTTTTCTTT GTCAATGATTGAAAATTTTCGGTTGTTAGGTTTTGAAATGGAATCCATTCCGTTATG[A/ ]TATATTAATTTTGTTAAATTGTAAGACAATATCTCTCTCAGCCTTCCACAAGACATAGTAGC CAATGTTGAATTTTCTTTATTAAGTGGATGTGGTTTTGTTGACAACATTTACTAAAAATC AATATGGTTTAGTTGATGGTTATTTTACAATAACATGTTTGTATGCTCTGATG
>Gm16_3 0539518	CAACAGGCGGACGGGCAAGGTTAACCGAAGGGTGCTCTTTTAGGAAGAAGGGAGATTGA ATGGTTTGTTAACAGACTTTTCTTTGTCAATGATTGAAAATTTTCGGTTGTTAGGTTTTTG AAATGGAATCCATTTCCGTTATGGTATATTAATTTTGTTAAATTGTAAGACAATATCT[C/T]T CTCAGCCTTCCACAAGACATAGTAGCCAATGTTGAATTTTCTTTATTAAGTGGATGTGG TTTTGTTGACAACATTTACTAAAAATCAATATGGTTTAGTTGATGGTTATTTTACAATAAC ATGTTTTGTATGCTCTGATGACGTGTTAGTGTCAATAATGTACATATGAATTTTG
>Gm16_3 0539519	AACAGGCGGACGGGCAAGGTTAACCGAAGGGTGCTCTTTTAGGAAGAAGGGAGATTGAA TGGTTTGTTAACAGACTTTTCTTTGTCAATGATTGAAAATTTTCGGTTGTTAGGTTTTGA AATGGAATCCATTTCCGTTATGGTATATTAATTTTGTTAAATTGTAAGACAATATCTC[A/T]C TCAGCCTTCCACAAGACATAGTAGCCAATGTTGAATTTTCTTTATTAAGTGGATGTGGTT TTGTTGACAACATTTACTAAAAATCAATATGGTTTAGTTGATGGTTATTTTACAATAACAT GTTTTGTATGCTCTGATGACGTGTTAGTGTCAATAATGTACATATGAATTTTGA
>Gm16_3 0626524	TATAATGAGGATGGTGTGAGGGAAGAGGTACTATGGTGAAGACTGTGATGTGAGTGATGT ACAGGAAGCAAGGCAATTTAGAGAGATCATTAAAGAGTTGGTGACGTTAGGAGGGGGCTAA TAACCCTGGGGACTTCTTGGCTTTGCTTAGGTGGTTTGATTTTGATGGTTTGGAGAAGA[A/ T]ACTAAAGAGGATTAGTAAGAGAACCGATGCGTTTTTACAAGGACTCATTGATCAGCATC GTAATGGAAAGCATCGTGCCAATACTATGATAGATCATCTTCTTGCTCAACAGCAATCACA ACCTGAATATTACACCGATCAAATCATCAAAGGACTTGCCTGGTAATTAATTAATTATT
>Gm16_3 1191863	CAACCCCAATCCAGTGATGCAGCTCATCTCCCCTGCATCCTCCACTGGTGAGAACACGCA GCACAACGCTGCCAACACCTCCAAGGCATCGGATTCCGAGAATTTCCGCTAGTCGGT GATCAAGGCTCCTAAGCAGGCCTCTGGGGAGCACAAGAAGAAAAAGAAGATCAAAGTGA[ A/G]GTTCCCATCAGGTCAAGAGCGGAATGCACCATCACAGGCAATTAGGAAATGCTTGCA CTGTGAGATAACCAAGACACCACAGTGGAGGGCAGGGCCAATGGGGCCGAAAACACTCT GCAATGCTTGTTGGCGTGCGCTACAAGTCAGGCCGGCTTTTCCCCGAATATCGCCCTGCA GCGAG
>Gm16_3 1204160	ATGTACTACCAAGCAAGGGGTCACATGCCACCTAAGTTTCAATTTTCAATTTTCCAGGAAT CAAATTGTTAGTGATTGATATTTTGTAGTGGTCAAAATGAATATTTAAAAATAAAAAATA AAAGTGAAGGAAAGCATGAGATAGATAATAGAGGAAACAACGTGAGAGTTCCA[G/A]JAG GAGTAAAGTAAAAGCTTTTGTAGATTCAGAAACCCCTCACTTGTGTTGGTGGTTGTGTT TTTGCATGCTTCGGACATGTCGATGGCGATGGTCCACCCTTCGGTTCAACCAATCGAAGC CCCACCCATGACTGAGCATGCAATTGCAATTGCAATGCCAGACACACTCTGAAGGA
>Gm16_3 1204451	CCCAATCGAAGCCCCACCCATGACTGAGCATGCAATTGCAATTGCAATGCCAGACACAC TCTGAAGGATACGCAGGGCATGCCGGCACCCCTCGGTGGCTTCTCTTGCGCTTCGCCC AGTTCTCTTTCGCCCTCGTTTCCCTCTCCGTATGGCTACCACCTCCGATTTCCCTTCTG[T /C]CACTGCCTTCCGGTATTCATTAATTCCCCCTTTCTTATTTTCTGTTAATCGTGCTTA TTTTAATTGGGTTTGTGGTAATTGTTGATTTGGGCTTGTTTTTGTGTTCAAAATCAGCAC CCCAGATTTTATTTTTTTCAAGTGGGTTTTATCAAGGGAGCAATGAAATGAACTCTT
>Gm16_3 1225684	TAACAAAACAGATCATATATCTGCCATTATTTCTAAAACAGAAAAGGATAGTTAAAATAAAGA ATACAAGGTTACTCCAGAATAACGATGTAGTAAAAGGAAAGTATAATGTACAAAGGTTTCT AAACAATACAATATGGTGAACAAATAGGATACCTGCATAATTTCAAGGTGCATCAG[C/T]CC

	CACACATTGAGGCTCCGAGAACCTTATCTGTCTCAGCATCAACAACAAGCTTCATAACAGT TTTTCTTGTGCGCTATTTGAAGAATATCAGCAACCATAATCTAGATAAGAAAGTTATGCTG CAAAGAACATAACTAATACTGGGTCCAACCTGAGATTGAAGAGCATGTGATGATAA
>Gm16_3 1229414	ACAAAGTGATTGATTTTGTATTTATCATTGTACCCAACTTACCTATTATAAATAGCATTTCAT GTTTGTGCCTCCAAGACACAACGTTTCAGATATTTCTTCTCTATAGTATCCTTACCAGTGGC AAACAGTACAGCATCCGCAATCAGCTCTTCACCATGATCTGTAATAACTTTAATGC[A/G]GT CCTCTGTTTTGATCAACTAAAATAATGAAATTTGTTACCATTAGCTAAGCCTCCAACCAAAG CAAATGAAAAAGTTCTACAAATGTGAGGTTGCAAACCCCTAAACCCAAAGGAAATACCTGT GTCAAATTGGTCCTTGGATGCAAATTAATCCCCTGCCTTCAAGATTTCTTGCCA
>Gm16_3 1461554	TGCAGAAAATGAAGGAAGATGTCATGCTGTTGCAACCTGCTTAACCAGGAAATTATCTCTT CCACCCTCTACATTTCACTCTTTTCTTGCACCATCATCATCCTCCAAAGTGACCCCAAC CTACTGGTTCAAGTAACAACCTTGATACCTGGTACACCGCGGCTTGACGAAGCCGTG[G/A] CGTGACAAGGGACAGAGTAAGAACTGGAATTTTGTGAAATTGCAATGGAGTCCTAGTC TAAACTATAAGATAGAATAAGACTAGGGAATTTCTTGTGGGATAGTTTATAGCATGTATT GCTATTTTGCCCCCTTTGAATTTTGTATGGGTGCAATTTATCTCTTTCTCTAGTTTTCT
>Gm16_3 1475163	TATAAGTGTTGTTTTCAAAATATTAAGGAGTTGTGGGGTGCGAGCGTGATTTTCTCGTTATA AAAATAAGAAGGAAGAGTGATTCAAATCTGAAGCTGAACCCTAACTAGCTGAGCAGACGG AGGAGAAGTCTGACTGAAGCTGCCTCTGCGACGCCGTTTCGATTACGCTCAGACATTT[T/ C]TGTAAGTTCATCGTTTACACTTTACACAAATACCCTATTCTTCGCATACCTCGTCGCTG CAACCCCGAACAACGATAACCTAATCAATTTGGCTTCGCCCCCTCTTTTTCTCTTACAAA CAATAACAATTCACGTGTGATTGATTTCTCCAGTGTTTTACGAGTTTCAAAGATGCTCG
>Gm16_3 1476359	CATTTATCAAAAGCCCAGGTTAGCAATGCAAAACAGTGAAAGTGCTGCTTGCTGCTGTAG GTTGCAGTCATGGTGCAAGAAAATGGCAAGAGGAAGAAACCAAACATTCTGGTGACTGGC ACACCGGGGACAGGAAGACAACCTGTGTGCACTGCTCTAGCTGAAGCCACCCAGCTCT[A/ T]TCACATCAATGTGCGAGAATTAAGTCAAAGAAAAGAACTTGCAATGATGGATGGATGATG AGCTTGATTGTTACCTTCTTAATGAAGACTGGTAAGCTATCAATTTATATAATTTGGATCA TTCCTATATGCTTGTTGAGATTAGCTCCTCTAAAGTAAATGAGGAGATGTGTAA
>Gm16_3 1516052	TCGTAAATTGCAATATGGGGATCTTTTTGGATTTTACTTGTTTTATTATACCTGACTTTTAG TCAATCGTTCAGCAATAGGTATATTAACCTTGAACGCTACATGGACAATCATTTTACTGTT TTGTTTATCCTAATAAAAAAATTGTACTTTTCATCAATGACATGTAGGAAAGC[C/T]GACAG GCGAGTCCTGACAAGAACTCAAATACTGTACATGGAGGAGTTGTCGTTAGTTTTAAATCAAG TTGAATCCATTCAAGACATTCTTCCTATAATTTCTGTCAATTCTTGAGGTAAATCTAAGCATT AGCATTTTATTTAAGAAATGTGGCTGTTACCCATTTTTTCATTAACCTGTA
>Gm16_3 1516059	TTGCAATATGGGGATCTTTTTTGGATTTTACTTGTTTTATTATACCTGACTTTTAGTCAATCG TTCAGCAATAGGTATATTAACCTTGAACGCTACATGGACAATCATTTTACTGTTTTGTTTA TCCTAATAAAAAAATTGTACTTTTCATCAATGACATGTAGGAAAGCAGACAGG[G/T]GAGT CCTGACAAGAATCAAATACTGTACATGGAGGAGTTGTCGTTAGTTTTAAATCAAGTTGAAT CCATTCAAGACATTCTTCCTATAATTTCTGTCAATTCTTGAGGTAAATCTAAGCATTAGCATT TTATTTAAGAAATGTGGCTGTTACCCATTTTTTCATTAACCTGTATAAGTCA
>Gm16_3 1827645	GCATGTGTTGCAACTTGTTTACTAGCAACTCTTCTGTATAAATACCATATACACACAATGA AATGAAAAGCTTGAGTTTCTTTCTTTCAATTATTCTAACAGAGGGTCGATCTAAAGTATTTA TGGAAGATCATGATTAGGGCATTATTTTCTTAATAAGCAAGAGAAAAATGGACA[A/C]AAT AGAAGCACATAATACTGAACACTTTATCCTTATTACATGAGTCATATGTCAAACAGGAAAC AACAATCCCCAAAATAGAACACACCCATATCCTTAGCAATTAACAACACCACACAAGAAG AACTACACATAACTTACAACTTTTGCTCCATTCAAACTGATCAACCAATTGCA
>Gm16_3 1827884	GAAACAACAATCCCCAAAATAGAACACACCCATATCCTTAGCAATTAACAACACCACACA AGAAGAAGTACACATAACTTACAACTTTTGCTCCATTCAAACTGATCAACCAATTGCAC CACCTCTTCCACCACAGACGTCAACAAAGACTTTCCTAGCATTACCACATTTGACTTT[A/C]C CACGAATTTACCTTGATTTCAGTGAGCATCCAATTGACTCAACTTTATAGTAGCATC CACAACTTCTCAAAGAACAACGTTTGGTTCAACGCAAATGCATTGACCAACCCCTTAGTC CTCTTATCATTGAGCAAGTCCTGGTCAGAGGTGAACACCCCTTGCGGTTTCATTAGG
>Gm16_3 1827991	ATCAACCAATTGCACCACCTCTTCCACCACAGACGTCAACAAAGACTTCTAGCATTACC ACATTGCACTTGCCACGAATTTACCTTGATTTCCAGTGAGCACATCCAATTGACTCAACT TTATAGTAGCATCCACAACTTCTCAAAGAACAACGTTTGGTTCAACGCAAATGCAT[T/A]G ACCAACCCCTTAGTCCTTATCATTGAGCAAGTCCTGGTCAGAGGTGAACACCCCTTGG CGGTTCAATTAGGTCAAGGTAGTATTTGTTGTGCAACACAGTCGGGGTTCTGATGTCCAAG TTGACGGTGTTGCCGGAGTTGCGTCGGGGCAAGTGGAATTGATTTGCTTGGCTAGGGT

>Gm16_3 1828137	CAAAGAACAACGTTTGGTTCAACGCAAATGCATTGACCAACCCCTTAGTCTCTTATCATT GAGCAAGTCCTGGTCAGAGGTGAACACCCCTTGGCGGTTTCATTAGGTCAAGGTAGTATTT GTTGTGCGAACACAGTCGGGGTTCTGATGTCCAAGTTGACGGTGTTGCCGGAGTTTGCG[C/ T]CGGGGCAAGTGGATTGTAGTTGCTTGGCTAGGGTTTTGTCCATGTTGGGGTCGAGAGG GGAGAGCCTGTTGAAGAATGTGCCGCAATGGGCACGACCAAAGGTGTGTGCGCCGGATA AGGCGACAACGTCGGTGACATCAAAGTTTTTGGCGGCGAAAGCGTCAAGGGTCACGCCG GTGG
>Gm16_3 1829738	ATATCACAAAGAAAAGCTTCAAACTATAGTTAGTAAGAGCAGAGGGTTTTGAGAACCAAA ATAGTAGTGCAAGTAAAGGTAAGAGGGATGAAGAGATTACAAGGAAAACCTGAATCACGTG CTGCCAAAACAGTGATATCTGCGCAAGAAACGATCCTTCCACACTCCTTATGGATAAT[C/T] GCTCTGATGTCATCAATGGTTTGAAGGCCCTCAGTTCTGATGCCTCCGTTGGCTGGTTGG TCTCTTTCACTTGGACTCCCCTCTAGCAGCAATGATCCATCACATCCCTATACAAATGCAA ATTAAGTTAGAGGTAATATCTTTTTTCTATATATATATGCATGCATATATTAACC
>Gm16_3 1840753	GTTGGGGGACTCGGCATTGGGACACGTAGCTATCAAATTGTTGTTGAAGTTTGGGTGCGAT TGGTGGGTCCGTTTTGATGGTTCTGTTCACTAGGGAGGGACAGTGGGCACGACCGTAGG TGTGTGCGCCGGAGAGAGCGACCACATCGGTGGCATCGAAACCTCTATTTCCGAATCCTC [C/T]CAGAAGATCGTCGGTTCGGAAGAATGGTGCCGGTAGGTTGTCCGGCGCGGTGGCG TTTGGTCTAGGCCGTCTTTTCTTCCAGTGGCACGTCAAATCAGGGCCTCCTAACTGTT CAAATTACAAAGAAGTTACCAAAAGGAACAAACGAGAGATCAAGCATTATGTTTAAATGTG TTT
>Gm16_3 1840819	GTCCGTTTTGATGGTTCTGTTCACTAGGGAGGGACAGTGGGCACGACCGTAGGTGTGTG CGCCGGAGAGAGCGACCACATCGGTGGCATCGAAACCTCTATTTCCGAATCCTCTCAGAA GATCGTCGGTTCCGAAGAATGGTGCCGGTAGGTTGTCCGGCGCGGTGGCGTTTTGGTCCT A[C/T]GCCGTCTTTTCTTCCAGTGGCACGTCAAATCAGGGCCTCCTAACTGTTCAAATTA CAAAGAAGTTACCAAAAGGAACAAACGAGAGATCAAGCATTATGTTTAAATGTGTTTTAA TCTCATTATAATGTTATAATTTGAGTGAAAATTATCTCTAATTTATATATTATAAATGATTT
>Gm16_3 1842815	GAGCTACTCCACTGTCCTTCTCGAAGACATCCTCAAGATGCTTCCTTATAATCCTCTCAAG CTTGGGACATGTCAACAAATAGTAATTCATGATAGTCCAGGGACTAGTTTCCAGAGGA AACATGGATTTGAGAAGCCAATACAAGGGAAGAAATCAAGAGCAAAGAGTGAAACAA[T/ A]GACTAACCCTAGCCATGTTTGATTAATTTGCTATGAGTGTGACTTGAGATAACCAAACT TAGAAGAGTTATTTATAGATTCAATATTGAGGTAACGTTCTTCGTTGCTTCCACTTCCCA CCCACTAGTCTTCTACGTATAATATTTAAACGTGGCGCAAGCGTTTTTTGAGATTTG
>Gm16_3 6812185	GTCCCTCCATGATAGCCCTCCATGTAGCTTCCAGTGTGTCATCACCATCTGAATCCGGCA GGACATAATCATTGGTGCAACTTTCAAGTGAAGGTTGGATCATGAATTTGTCATAGGTCAA ACCAGACTCTTCAATCACCTCTTCTTGTCTAGCTCTTCTTCAACCACCTCTTCTTGT[G/A]C TAGTTCTTCTTCTTCTTCTTCAACCTCCTTTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT TGACTCAGTTGGGTGGTTGGCAGTGTCTGAAATGACGGTGGTTGTGTGCTTAGGATCAGA GGGTGCTGGTGAGTCAGAGAAGGTTTTGGGGGGGAAGTTGGAGGAAGCTGCAATGG
>Gm16_3 6812187	CCCTCCATGATAGCCCTCCATGTAGCTTCCAGTGTGTCATCACCATCTGAATCCGGCAGG ACATAATCATTGGTGCAACTTTCAAGTGAAGGTTGGATCATGAATTTGTCATAGGTCAAAC CAGACTCTTCAATCACCTCTTCTTGTCTAGCTCTTCTTCAACCACCTCTTCTTGTTC[G/T]A GTTCTTCTTCTTCTTCTTCAACCTCCTTTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCT ACTCAGTTGGGTGGTTGGCAGTGTCTGAAATGACGGTGGTTGTGTGCTTAGGATCAGAGG GTGCTGGTGAGTCAGAGAAGGTTTTGGGGGGGAAGTTGGAGGAAGCTGCAATGGTG
>Gm17_3 054922	AACGCTCATAGGTTTCATTAGTAGTTTGCAGAAGGTAAATTGCAGTATTCTATTAACCTCAATC TCTACTCCATCTATAGTATTCTCAAATCTGAATTAGTACATGTTATTTAGACATTATATGAA TATAGGTCTGAAAAGTTGAAAACAATATGCAGGGTTATGACACGTTAGTAGGGG[A/C]ACG AGGAGTTCAATTATCTGGGGGACAGAAGCAGCGTGTGGCAATTGCACGAGCTATAGTGAA GAACCCAAAAATATTACTGCTAGATGAAGCCACAAGTGCAGTGGATGCTGAGTCTGAAAA AGTGGTTCAGGATGCACTAGACCGTGTCTAGGTGGACCGAACCACAATAGTAGTGGC
>Gm17_4 444643	AACTGGAGCAGTGGCTTCAGTCTTCTTGGTATGTATTTTTATGTTTCCTTTGTTTCAGACTT TTTTTACATGAAACTGGTATGATAAATGATATGGTAAACTTATATTTATGTGTCTATCTTCT TGGCAGGGTTTTGGGTCCCTGTTCTCCTACTTGCTTCTGGCGTTTATGTCTAA[T/C]ATAA ACATTGAATTGGATGAAATTTTGCCAAGCAGATAGCTTTAAATGAAAAATTGCATTGGACG TTTTAGTTTGATATCTGACTCTAAGTTATGTTTCATAATCTATCGATGACAATGTTGGTTGTG TGTAATGTTCAAGAAGAATAGAAGATTATGATATCTGAGATAGTTTTCAAACA
>Gm18_1	ATCGGATTCAGGTGAAGAAGAAGAAGAAGAAGAAGAATATTCATCGGAGAAGGCCGAGG

330362	ACGATGATGTTTGGTAATTCTCAACCTTATTTAAATCCTGAAACTCCGATGTCTCTGTCTCT GCCTTGTGTATCTAGAACAAATAGGGTTTTGAGTTTTGAGTTTTCTGCGAATTGGTCC[A/C] CAAGCTTTGACAAATGGCTTGAAGCTTCGCATCGACCGCAGAGGGGAGAGGCTTGTAGTT CCCAAACCTACCGTACGCGCGTGCTCCATGGTGACGCAGGTGAGGGAAGTTGAGCCTCG CGTTTTCTCCTCTTAGCTTAAACGCTGCGTTGTCGTAAGCCAACGCTGCTTCCTCTGCGGT
>Gm18_1 330743	ACGTGTTTCGGTTCTTGGGGAGTCTGATCTCAGCGACCCACTTGCCCCAGTGTCTCTGCCT CACCCACGGTAAAGCTTCGTGGGCTTTGAAGAAGGAGAAGCCCGCACGTGCTTCATGG GAATGGGCTTTGGGCTCAGAACTGGCCTCGTGGAACCTGGATTCTGGGCTTGAATTTGAA [T/C]CATCTGTGATGGGGTGAGTTGGTTTCAGTTTGATGGAAGTAGCATGGTTTGGAGTTAT TGAAACCTTTTGCTCGTAGCTTGTGTTGAAGAGAAATTGGGAGAAGAAGAAAAGGAAGA GAATGAAGGGTATGTAATGCTAGATGAATCACTTTTCATAAAGGCTCAAGTGCTTTCATC AG
>Gm18_1 330746	TGTTTCGGTTCTTGGGGAGTCTGATCTCAGCGACCCACTTGCCCCAGTGTCTCTGCCTCAC CCCACGGTAAAGCTTCGTGGGCTTTGAAGAAGGAGAAGCCCGCACGTGCTTCATGGGAA TGGGCTTTGGGCTCAGAACTGGCCTCGTGGAACCTGGATTCTGGGCTTGAATTTGAAACA [G/A]CTGTGATGGGGTGAGTTGGTTTCAGTTTGATGGAAGTAGCATGGTTTGGAGTTATTGA AACCTTTTGCTCGTAGCTTGTGTTGAAGAGAAATTGGGAGAAGAAGAAAAGGAAGAGAA TGAAGGGTATGTAATGCTAGATGAATCACTTTTCATAAAGGCTCAAGTGCTTTCATCAGC TC
>Gm18_5 842068	TGTCTTATCCTGCAACCCAGAAAAGAAGGGTTAAATTTCTTTAGCCTCCAAAACCGTAGAGG GTCCTTCCCTGCCTCTTGAGCGCATAAACACATCCATGGCAGTCACCGTCTTCCTCCTA GCGTGCTCGGTGTAGGTCACAGCATCTCGAATCACGTTCTCCAAGAAAATCTTCAGAA[C/ T]CCCCCTGGTTTCTCGTAGATCAAACCACTGATTCTCTTCACACCACCTCTTCTCGCTAA CCTACGAATCGCAGGTTTCGTGATACCCCTGAATGTTATCGCGAAGAAGTTTCTGTGCCT CTTGGCGCCTCCCTTGCCCAAACCTTGGCCACCTTTTCCACGACACAGACATTTTCTCGAG
>Gm18_4 8861899	CATAACAATTGTATCTTACTGATTTTTCTCATTTTCATCGTATCTAAAATCTTAAATCTTCTAT AATTGAGACATACACCATATCATGTAAAACAAAAGAAAATTTTCAAGTGAGAAAAGATAA GTAAAGAAAGTATATATATCACAGCTCAGTATAATGCAGAATCACCTAGTTTCCC[A/T]TCT GCCATGGAGAAGGGAATGAGGTGAATTAGCAAAGGAAAGAAAAGTAGACTCCATGTCTCT ATTTCCAAAATTGGAGCCTGAACCTCATCTTATCAAAGGATGAAGATGCAAGTGAGTGCAAG AACTCATCAAACCCCTCATGGTGACTTATATCCCCTGGCTTTTTCAAGTCCCTCTGGG
>Gm18_4 8861902	AACAATTGTATCTTACTGATTTTTCTCATTTTCATCGTATCTAAAATCTTAAATCTTCTATAAT TGAGACATACACCATATCATGTAAAACAAAAGAAAATTTTCAAGTGAGAAAAGATAAGTA AAGAAAGTATATATATCACAGCTCAGTATAATGCAGAATCACCTAGTTTCCCCTC[A/T]GCC ATGGAGAAGGGAATGAGGTGAATTAGCAAAGGAAAGAAAAGTAGACTCCATGTCTCTATT TCCAAAATTGGAGCCTGAACCTCATCTTATCAAAGGATGAAGATGCAAGTGAGTGCAAGAA CTCATCAAACCCCTCATGGTGACTTATATCCCCTGGCTTTTTCAAGTCCCTCTGGCAC
>Gm18_5 9613520	GTGGGAGCTGGAGATGGTGGGAAGCTGTGAGTCCAAGTGGTGAACGTGGAGGCAGTCTT CTACGGAGCCATCGCACAGTACTGAAGTTGTGGTGTGGTTGGACTTGGACAACACTCCTG AAATGCTCAAATGTACCATAACCCACATCAGCAAACGCCTTGATGCTAATTGCTTCTTCA[A /G]GTCTGCTAGCTACAAAGTCTTGAGAGATACTACTATTTTCATAAATTGCACACTGCATG TATAATATCCACCCTTGATATTGATATATGACTACCAAGGTTCCGCTACATTGAGTGCC AGTTAATTAAGCAAATTATTATATGTGAAATGATAATCATTGTCCGAACATTGAATAAAG
>Gm19_3 81220	GTTATACAAGTGACATAATATCAACTGTTCAATTTCTCATCAAAAACAAGGGGATTACTTA CTTTAGTCTCTACAACGTGTTTTCACTTTATAAGGGCCAGATCCAGCATTACAGATTTACAT AAGTTTGTTAATTTAATCTACAATTGATGCACTTATTGGTTTGGTGATAATGAG[A/C]GTGC AGTGGTATGGAATCCCTGGGGTAAGAAGGCCAAGGCTATCCCTGATTGGGAGATGATG ATTACAAAGATCATGATGTGTGTGAATTCTGCAGCTATCGATACCCCTATTCTCTTGAAGCC CTCTGAAGAGTGATGGGATATCAGGAAGTCTCTACGGTTTCATCAAGCTATTGCA
>Gm19_1 210520	TTGCATTCCACTATGTGATTATAGGTGCTCATTACACTCAAGGAAGAAATTGTGCATGAGA GCATGCATAACCTGTTGTGACCGATGCAAATGTGTCCCTCCTGGAAGTTATGGTAACAGG GAAAAGTGTGGCAAGTGCTACACTGACATGCTGACTCACGGCAACAAATCAAGTGCC[T/ C]ATAGAAGAAGCCTAATATCTAGTAACCTAACCTAAGCTTTTTTGTAAATCAAGTTTGAATCA TGAGTAATGTGGTTTGAAGTTGCTAGTGATTTAATAACCGAGAGTGATAATCATAATTGTA CAAGCTATCGTGTTAATCAAATAGTCAACACTGTTTGTGTTGTCTATAGGATCCATT
>Gm19_3 750352	TCAAGTGACAGCGTGGTGGGTTGTGATTCCAATCGAAAAGCCTTGGTTGGGGAAGAGGGT GAGTTGAAGTGCTAACACGGAAGCTTCTTCATGGGAAATGGTTAAGTGGGGTATTAGGTT

	TTGGAGGTGGTGGTTGACTTGAGAGAGGTTTGAAGAGAGGGTGGTGAATCTGCGTTGGA [G/A]TGGGCGATTGTGAAGGGGATGGTGTGCCGGGAGTGTAGGTTATGAGAGGGAGGG GTGAGTGATTGGGCCATGTGATGGTGCCGGCGAGAGGAGGGAAGTGGTGGAGAGTGAG GGAGAGGGAGTGTGGAGATTTGGGAGAACGGTGGTGTGCGAAAAATGATGTTGTTGG GTTTGGG
>Gm19_3 4933973	TTCGCACAGAGCGGTGAACCAAATTACATAGAGAATTGAAAGGAAAGCGAAGAGATCGAG GAGCGAACCTGGGGTATCGGAGATCTGGAGAGATCCGAAAGCGAGGTTGATTGGTGAGG GGGAAAGAGAAGAGAAGGGTTGAGGAATAGGGTTCCGGTTCCATGTTGGGGATTAGGG T[C/T]GGGAGATGGCGCAGAGGAAGAAGACGAAGATGCCGCTTGGGTTTGTGAGGGCAA GATGGACATGAAAGTTCTAGAAGGAGGGGTTAGGGTTGTTTCGTAGGTTTCAGCTTCTAT TCTACTTAACCAAATCTATGTCGAATGTACTTATTGTGCTGGCTGTGTCTTCGCGGCTCCGT GGTA
>Gm19_4 0301455	AAATGAAAATGTAATACTAATCGTTAGTTGTACGAACCACCAATAAGCAGGTTGGTTGAGA GTAATGTTAGAGACTTGAACGTTGTTGCAATCGAGAAAACCGATGAGCCTGGGTCTGCAT TCGTCTCCCAAGCAAGCCCTGTTGATTCCAACTAACCTAACGTTCTTCTCGGAT[A/T] TTCCCTCACCACGAACCTTCGAGCTTGCCCGTCCACCGCTCCGCCGCCGCGGATTCCCA CGTCCGTCGCGTTCTCCGCAACAACCACGTACCACCGCCACGACTCCTCGGGTAGTCC TCCAGCCTCGTGCCGCCAAGGATGGTGGCCCCGACTCCACGTTGAGCACCACGCCGGA TTT
>Gm19_4 2879589	TTGAAAAACCCCTTACCTTCTTCAAGTACTTGTCTACAAGCCAAGGCACCTGTGACCTGCT CTTGAAGCCACCAAAAGCTGTTCCCTTCCAGACTCGTCCACTCACCACCTGGAAAGGACG GGTTGATATTTCTGCCCTGATGCTGCAACACCTACAATAACTGACGTCCCCCAGCCC[A/T] JGGAAATTATCAAAGAATATTTTCATGCCTAGTGTAATAATGTTCAAAAATTGTGTCAATATAA CAGCAGCAGCATTTACCTTGTGGCAGCATTCCAACGCAGCTCTCATCACGGAGACATTTT CAATACACTCAAAGCTATAATCAACTCCACCATCTGTGCGATCAATTATGACCTGCTG
>Gm19_4 8261364	CTTGAAGAAAGGCCGTTACTCTCGCCGCTTGGCACCGGCGCTCCCATTTATCTCGCCGC CGTTCTCGAATATCTCGCCGCAGAAGTAATGATAATTTACGATTTCGATTATCCTCTTCT CGTTAACAGTTAAATTATATCAAATTAATCAATGCTTACGGTGGAATTTTCGAAATGC[A/G] GGTGTGGAGCTGGCCGGGAACGCTGCACGTGACAACAAGAAGAATAGGATTAACCCTA GGCACGTGTTGCTGGCGGTGAGGAATGATGATGAGCTGGGGAAACTGCTTCAAGGTGTG ACTATTGCTAGCGGTGGAGTGTTGCCGAACATTAACCCGGTTTTGTTGCCTAAGAAGAGT AC
>Gm19_4 8638348	ATTTGGTTTGTGATGATGTGTGTTTTGTTAGGTTTTGGAATTGGCTGGGAATGCTGCT AGGGATAACAAGAAGAATCGTATTGTTCTAGGCACATTACAGCTTGCTGTGAGGAACGAT GAGGAGCTGAGCAAGCTTTTGGGATCTGTTACCATTGCCAATGGAGGTGCTTGCCGA[G/ A]TATTATCAGACTCTGCTTCCCAAGAAGGTTGGCAAGGGGAAGGGCGAAATGGATCA GCTTCTCAAGAGTTTTAGGGTTTTGTTAGGCCTATTTGGACTATAGTTTTTATTTTATTGC GGTTGTAATATACATGTTTTGGTTAATTTTTAGATATCTGCTCTGAGGAATTAAGTGTTT
>Gm20_3 9267808	TTGCCCCACCGTTGCCTCCAAAGATAAGCAACTTTCATCCATAAATACCTTTCATTGTTAG CGTGTGTACTCCACATCCTCCATTAGTTACCCCTTCTCAAGCACACGTTACCCCTTTTTT TCATGGCATCAGATCACTCGTCCGTCAACCAACCCAAATTTGTCTATCCAAACAATG[A/C]A CTTGAGCAACAAGAAGCCCTAGAGGTGCAAAACCAAGGACTTTGACTACTCAAAGAGATCT CAATGGCTACGTGCTGCTGTTTTAGGTGCCAATGACGGGTTGGTCTCAACAGCATCACTA ATGATGGGTGTGGGAGCAGTGAAGCAAGACATCAAAGTCATGATCCTAACAGGTTTTGC
>Gm20_3 9367321	AGTAGTAGGAGTAGCAGCAGCAACAACACTGTGACCATTATTCATTTTCTGAATTTCAATG TTGTTATTATTTGACATCATCAGTGGTTGTGGAGCTTGTGTGGTGGTGGTGTACTTGT GTGGAAGTGTTGTTGAAGGAAATTGGCACCAGCTTTTGTGTGGTGGAATTTTGGT[T/C] TTGACCAGATAACTGTTGCAAAAAAGCAATAACTGCTGCATTTTTTGTCTGCTGCGGTTGAT CTTCTTGGACAAGAAGTTCGTGTTCTCTATTGATCCTTGCCATTTCTTGATCCTCCAAG CCTCTTGTGCGCCACTTGTTCTCTCTCCCTATTGTCAATGGCTTCGAGGAACCTCTT
>Gm20_3 9367375	TTCAATGTTGTTATTATTTGACATCATCAGTGGTTGTGGAGCTTGTGTGGTGGTGGTGTG ACTTGTGTGGAAGTGTTGTTGAAGGAAATTGGCACCAGCTTTTGTGTGGTGGAAATTTT GGTGTGACCAGATAACTGTTGCAAAAAAGCAATAACTGCTGCATTTTTTGTCTGCTG[G/A] GGTTGATCTTTCTTGACAAGAAGTTCGTGTTCTCTATTGATCCTTGCCATTTCTTGATC CTCCAAGCCTCTTGTGCGCCACTTGTTCTCTCTCCCTATTGTCAATGGCTTCGAGGAACC TCTTCTGATTTCTCTTGTGCTTGGCCAAGACTTGCTCGTGAGCCTCCTGAAGTAGTC
>Gm20_4	TAAAAATAAATAAATAACAAAGTTTATGCTAAATTACTTTGAAGCCCAACAATAAAAAAGAGA



1862066	ATATTATTAACCCGAGGTATATAAACATATCCGAATGAGGACGCACACTAAACTGGGGGG ATATGCTCGATTGAGGGTGACGAGGGATAGCAGATCTGAATCTCATCTTCCCCAAA[C/T] CCTCTCTTTCTCTGATCTCATTTTAACCCTAACCCTAACCCTAACCCTAACCCTAACCCTA TCAGAAGAAGATGTTGACGAAATTCGAGACGAAGAGTAACAGAGTGAAGGGACTGAGTTT CCATCCGAAGCGGCCATGGATCCTGGCGAGTCTCCACAGCGGCGTCATCCAGCTCTGG
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**Supplemental Table S3.** SNPs identified by Method 1 on the four-library comparison

This table contains 261 SNPs identified using Method 1 on the four-library RNA-Seq comparison between the Clark/IsoClark 19-day root and leaf experimental data. The file contains a unique SNP identifier that includes the genomic location. The sequence surrounding each SNP based on the soybean Glyma1.01 genome assembly is also provided.

SNP	[Clark/IsoClark]
>Gm01_55645316	GCCGATTAGTCGCCATCACTGGCCGTGTTTCAGAAAGGTCAGGGACGAATTGGAGCACCTGC TCGACGACGACGATGATATGGCTGAGATGTATCTCTCTGAAAAGCTAGCCGAACAAGATGA CTTGGAGAAGGCGGAGGATACCTCTTCTGCCGACGATGTTGATGATCACATTGACAGG[A/C ]ACCTCTTCCTCCCTTTTGA CTCTCCTTCTTCCCTTCTAACCTCTATATGTAATTCATCCTT CTATTTTTTTTTTTTTTTATTTAGGACCGCTCCTGAAATATCGTTGGACAATGTTGTTGGTAGG GACAGCCATGGAACTCGCGATAGCGCCACCTACAGCGCCGTCACCAAGCAACTT
>Gm02_42632633	GTTGGCATGGCCATCCACCAAGGTCATTCTATGTCCGCCACGGCCCATGATGAAGGGTT CCGTGCCATGTTGTTATTGGCCTCGCACGTGCCAGGGTAGCCTTTGCTGCTGCTGCTGCT GCTGCTCTGTGTGGCAGTGTGAGTACTGGGGGAACATGAAGAACTCAGAAAAGGGTG[ A/C]TGAGTCAGAGAAGCCACTACCCTCCTTTGTTCTCTTCTCTGCCCCTTCATGCACTGCA GAAGCTGCTCCAGCTCCTTCACGAAGTTAATGGTGCCACCAATAATAGAGGCTTGATCACC CTGTTTTTTTTTTTTTTTTTAATAACCGCACATGTGGCACGGTTAGTTGGCTAGCTAGTA
>Gm02_42829778	CAGCACCGACGGCGGACTGTGCCTCACGTGAGCGCGTGCCTCCCACTCGCGATCTCGA GCAGCAAACTCCGAACTGAAGACATCGCTCTTGAACTTAGATCTCCCGGCGCAAGATA GCACGGGTGCGAGGTATCCTAACGTCCCCGCCGGTGGCACGCGAGAATCCGCCACGTGTC[ T/A]CCTCACCGCGAGCCCGAAGTCACCGAGTCTAGCGTTCCACTTTTCGTCGATTAACACA TTGGACGATTTTATGTCCCTGTGAATAACTGGCGGTTTCGGAAGAGTGTA AAAAGGCGAACCG CTTTCGCGACCTGAACCGCAAACCGGACTCGCGCGGTCCAGCTTGGGGGTGCAACCGGTT T
>Gm02_42830291	TTCGATTTGTTTTGAGGTTTTGGGTTTTGGTTATTTTACGGCGGCGACGAGAGCGCCGCG TGGAAGGCGGCCTTG TAGACTGTCCGTGGCTGCCTTTACCTAGGAAGGTGTCGGCAGAG AAGTTGTTGGTGCGTCGAGGATGTGCGAGTAGGCGAAGTGTCGACGGGTGGGCCT[C /G]GGCTGGGCTTTTGGGCTTTTTCTTCTTCTTGGAGGAGTGAGGGTTCGCATGTGGCTATTG CGGACTCGGCGTTGCAAGTGAGGTAGGGCATTGTGAGTAATGATGGAGAAAACTAGAGAA GAGAGAGCATAAATACTCTGAACACTGAACAGATACATTTTGATTTAAAGGGAAATCAAT
>Gm03_36398914	AATGATATTTTAAGCATCTTAGGCTTTAGAATCATGTTTAAAGACTACTGTAGTCAAGTAAAG CTAATAACTCAAAATACATACATATTCCAGTGACAGAAAACAAGGATTATGTATGTAGTGTC CTGCCAAACCAGTTTCCAGTAAAAACAAGCTGCATGTTTCTTATTTCAAAGCT[A/C]ACAGG AAGGTGATAATCGTGACGTACACAGCATACAACCTTACATAAAAGTTTCTTATGGCATAGAAA ACATGTAGTCCTCCTATTTCGTACACAGTCAAGTTTAACTAAACAAACAAAGAACTCAAAGA TGAGAGTATAGGAAATACAAATACAATACTAGAGTCATTCTGTACAAAAGG
>Gm03_36398969	AAGTAAAGCTAATAACTCAAAATACATACATATTCCAGTGACAGAAAACAAGGATTATGTAT GTAGTGTCCTGCCAAACCAGTTTCCAGTAAAAACAAGCTGCATGTTTCTTATTTCAAAGCT CACAGGAAGGTGATAATCGTGACGTACACAGCATACAACCTTACATAAAAGTTTCT[T/A]ATG GCATAGAAAACATGTAGTCCTCCTATTTCGTACACAGTCAAGTTTAACTAAACAAACAAAGAA ACTCAAAGATGAGAGTATAGGAAATACAAATACAATACTAGAGTCATTCTGTACAAAAGGTT TGGGTTTCATCATGTAATAAGACAATACACATACTAAGACACATAATTCTCTCC
>Gm03_36399038	CCTGCCAAACCAGTTTCCAGTAAAAACAAGCTGCATGTTTCTTATTTCAAAGCTCACAGGA AGGTGATAATCGTGACGTACACAGCATACAACCTTACATAAAAGTTTCTTATGGCATAGAAAA CATGTAGTCCTCCTATTTCGTACACAGTCAAGTTTAACTAAACAAACAAAGAACT[C/G]AAAG ATGAGAGTATAGGAAATACAAATACAATACTAGAGTCATTCTGTACAAAAGGTTTGGGTTCA TCATGTAATAAGACAATACACATACTAAGACACATAATTCTCTCCCGCCATCTCTCCTCGT ACTTTCCAAATGAGTAGGTAGTTGAAGGCATCCCAGTCAAGTGAAAGGGGTC
>Gm03_3641699	GATGGAAGGTTCTTGCTTTTGTGTTGTATATATACATACTATTATTGCTAGGAAATTGAAGACC TAAGATACAATAGAGATGGCAAAGCTATTTTTGAAACAGGCAAAGCAATACGCAGATGCAA

2	GACCAAGCTATCCTCCACAACCTCTTCCAATTCATTGCTTCCAAGACTCCCTCTCA[G/A]AAC CTCGCTTGGGACGTCGGCACTGGGAGCGGCCAAGCTGCCAAATCTGTAAATTTCTTTCTT CTCCTTCTCTTTCTTAATTCCTTTCACCTATATATATATATATATATATATTTCTTGGTCAATTC CACAAGCACTCTCACATATTTTACACGGGCGTGAGATCCTGCTTAACCTAG
>Gm03_ 3641809 7	AAATGGGTTGATGACAATTATAGAAGCATTGATTTTCCATTTGAGCCCGTGGATGGAGCTG ATCACACAGGACCCTTTGAGTTTGTGACGGAACAATGATGGATTTGGATGATTTCTTGAC CTACATAAGATCATGGTCAGCATATCAGACGGCTAAGGAGAAAGGAGTGGAGCTTCT[A/C] GCGGAGGATGTGGTTGAAAAATTCAAGCTTGCTTGGGGTGAAGATGCTAAAAAAGTTGTCA AGTTTCCAATTTATTTGAGAATTGGAAGAACAGGGGATTCCTAAAGACATATGCAAATGGTT GCTTTTACTGTGTGGGAGATGTGACGAGTACCAACTTTTATGAGTTTATCCATTGAT
>Gm03_ 3646037 4	GACATTGCAAATTAATAATTTTCCATACAAGTTTCCTCAATCCTAATATAGTGATAGTGATAAA TAGAACTAAGACAACAAATTAGTACTAGGCTCTGTCAATCTTCATAGGCCAATACTTTTACC ACTCTTTAATAACTTATGACCCGATATTTGAAGGTGAGGATAACAATTGTAACA[A/C]GAAAA TTACAGGAAATACGGATTTTATTACATGCACGTAAGAAACACATCAAATTTCCACTCTCAAG AATCCAAAGAAAAGATAACAATGCATGCTCTCAAATCCTCAACATGCTAACTAACATCTTGC CAATGCAACTTCCTAAGGATGAATATTTAGACAAAAAACAATTAAGAG
>Gm03_ 3655394 4	TCTTCACGACTCATATGATACTCGTGACAAAAGACATACTCAGCAACAACCTTGAGGAACATT AACAACCTGGCTCCTTAATTTTGGTGACACCAGCAAGCACTGCAACAACCTTTCTGATCTTTGC CATGCTCTGTGAGCAAAGGCAGTGGTCTGCTCACCCTGTCTAGATGCACTATG[A/C]GC ATCAGTGACATCAATCGCTCTTTGATGTGAACCTCTCGTCCACTCTACAACCACCAATTATCA TAGGTGAAGTCATTTTGTACATGAAAGTTAAAAAATAAGATGTTAGGCCGATTTTATTTAATG TAATTGTGTGTAATAAAATACATATAAATGCAAATACATTTAATATACTTTTT
>Gm03_ 3655985 7	TGTTTCTGCAGGTGGAaaaaactCATAGATTAGAGTCCGAAATTCCTAAGTGAAGCTTTTG TCAATATATGAGAAGCAAAGGATTTTCTAATCAAACGATCTTAAGTGCTCTGATGTTAAATTT GACACCTAAAAGAGTCTTCACATATTTTGCATGCATGAATGGCAAGTTCCGAAA[A/T]TAGA ATGGAGTAATCACATCACATGCCGCTTCTATATACATATGATTTTGGTAAAAAAGCAAGTGGC CGGCAGGAAGGTGCATGGATTTTAGGAGGAGTTAAAGAAAAGAATATTGTATAAACATAAA TGTTACAATCAGCCTCAGTAGCAAAGGTGTGACACCCTCTTTTCTCACAATA
>Gm03_ 3655992 6	ATGAGAAGCAAAGGATTTTCTAATCAAACGATCTTAAGTGCTCTGATGTTAAATTTGACACC TAAAAGAGTCTTCACATATTTTGCATGCATGAATGGCAAGTTCCGAAATTAGAATGGAGTAA TCACATCACATGCCGCTTCTATATACATATGATTTGGTAAAAAAGCAAGTGGCCG[G/A]CAG GAAGGTGCATGGATTTTAGGAGGAGTTAAAGAAAAGAATATTGTATAAACATAAATGTTACA ATCAGCCTCAGTAGCAAAGGTGTGACACCCTCTTTTCTCACAATATATGTACTAATAATGA AAGGAATAGAAAGTTAGAATTAATGAAAAGTTTTAAACACATTTAATAAAAA
>Gm03_ 3656000 2	CATATTTGTCATGCATGAATGGCAAGTTCCGAAATTAGAATGGAGTAATCACATCACATGCC GCTTCTATATACATATGATTTGGTAAAAAAGCAAGTGGCCGGCAGGAAGGTGCATGGATTT TAGGAGGAGTTAAAGAAAAGAATATTGTATAAACATAAATGTTACAATCAGCCTCA[G/C]GTA GCAAAGTGTGACACCCTCTTTTCTCACAATATATGTACTAATAATGAAAGGAATAGAAAGT TAGAATTAATGAAAAGTTTTAAACACATTTAATAAAAGCATTTCAAAAAGGATAAAAGGTT TACATTCGTTTTCTAGCATTCTAATAAACTTATTTAATAAATAATAAAAA
>Gm03_ 3661293 8	GGAGGATCCGGATCCGAAAGAAGGCGGGTCCGAGCGTGAGACTAAAGATGGTTCCTTTTT ACGAAAGAAGCGTTTGGGGCTTAGTTTGAGGTTTTTGGGGTTGATGGTGCTAATGAGCTTC ATTGTTTTTGGAAAAGAAGAGAAGAGGAGAGTGGGGTTGAGGGGAAAGAGAGTGAAA[G/ A]GTGGAGGGATTAAATGGGGGCACCATGGAAGGTCCATTTCTCACTCCTTGCTACCTTCC ACGAAGTTTCTATACATCATGGACTCATTATAAAAGAAGGTGGCTATTTTATTTTTTATTT TTTTCTTTATTAATTAATATGTCATAATAGTGATTGTATTGGTTTGAATGCCTCAT
>Gm03_ 3663436 1	CAATATGGAGGGCAATAGGAACAAAATATTAATGTCACTGGAAAAGAACTGATACTTGA CACTAAGTGTCAACAATGCTTTGATCCATACCCGAATCTGTGCCCGAATGATGATCAATGG AATTGTAATCTGGCATGTGAGTTAAGGACCTGGCTGGCATGGATTCTTCATCATCTC[T/C]G TGTTTCAGATGCAGAGCTAGCAGCATCTATCTCCTTCCCTGGACACTCATCTTGGACACATT CAGCAACTTCAATGTTGTAATTTTCAAACCATGACTGCTACTACTTAAGGGTGACAAGTCT TCTTTCTTGACAACATTAACACAATCCCATTATGGATCCTTTCAGGCTGCAGAAG
>Gm03_ 3670959 0	GATGATATGTTTGCATCATGTTATCCGTAGCTTCAACTTTCTGTAACCTCTTGTATAGGCAAA GGTTTTCAAACCTTCAATCTATCCTTCTCTTGCTTTTGCCAAAATGTTATGGTGAATAGGGAA GGAAAAAGATGGAAAAAATGTGACTGTTTCATGTACTGATATTGTAAGGCTG[C/G]TAAA AGTATTCAAATTACAAAATGTAAAAAATGTAAAATGACTTCTCTGTTGAGATTTGTTTTCT TATGGCTTTCACCACCGTCCTGAACTTTAAATGATATTTGAAAAACAACATTAATTGGCG

	CATTCAACTTCGTAATTGCCTGGCGGCGTTCCGGCATTTTAACTTTTTCTTT
>Gm03_3680889 2	TGAGTCCCAAGGCTTCACTTCATTTCCACAGTAATGTACCTGAATCTGACATTCCAAAAGT TTCTGACATTGTCTTGCACCCATCTGTTTGCGTTGTCTTGGCTGGAAGCTACATATTGGAG GTCGATGTTAGGAAGATCTAGCAGCAACTCAATGTTGGAATCTCTCAGAGCTTGGAG[G/C]GA CTTCATGGTTTTCGATCATAGATTCTCATCCTTTGGAAGCCGTAAGTGTAAAAAGGGAAACT ACTTCTTGTGGCGATGGCAGATTGTTGCCAACTCTTCCATAGCATACACCAGATTGTGCAC CTGCACCAGATGGAACAAAATGGTGAATTGATAATACTACTTCTAAGATATCAAG
>Gm03_3695239 4	AAAAGAAATAAAAAAATAAGATCTGTTTTTTTTTTTTTTGAGAAAGATCATTTTTATTAT TATTGTTTTGAATTTGAGAATTATGTAATGATGAGAGGTTGATGTTGTGAAAAAATAATG GAAATCCAGGGAGGAGGCGGTGGTGGAGGTGGCAAGAGTGGAAGACAATGT[G/C]CACC AACAGCAAGGAGAATTTGTTCAAGTTTTCTTGTACACTTTGGTACTCTTCAAACATTGGT GTGATCCTTCTGAACAAGTATCTGCTCTCAAACATATGGATTCAAGTTCCCAATCTTCCTCAC AATGTGCCACATGTGCGCCTGTGCTGTCTCAGCTATGTCTCCATTGTGTT
>Gm03_3695995 5	AACGAATTGCAAAAGTTTTGGAAGTGTCTCAAATAAAAAATGCTCCTCAAACAGCTGGTTCT GACATAAACAGATATAAACTAGTTATGTTTCAGTGCGCAGAGAGGACTGTTGTTGGCATATTT GCCTTCAATCATTTCTTTACTCAACACCCATATATGTTTTGTCATGCCTGGAAT[G/A]TAGT ATGGTATCAAGTCTTCTCAAACACTACATAGAATTAGAATGCGGTAACGAATGGCAGCATTA AAGCACACACTACACTAGCAGAAGAAATCAGCAAAGCAAGCCGGGTAATGCCCAAATGTAA AAAAAGTAGCTTGGTTAGGCAATTCATGTCAATGTTTCATCTACTTGTTTTTCG
>Gm03_3696027 5	GTTAGGCAATTCATGTCAATGTTTCATCTACTTGTTTTTCGAAAGGTTAGTGTCTTAAAAACC AAAATACGAAACCAAAAAATGTTTCATGACCCAAAAACAAAAGAAAGGGAACGAAACCAT ACTCTTCTAATATTTATGCCAGCTCCGGACCAGATGGCAGTGCCTCATAACCATC[T/T]CG GTGCTGCTTACTAGCATCTCTTCAACTGGGATGCACCTGAGGATAGCAGCTATAGGCATG CTAAATGCTCCGATTACCACACTTAGTAACCAGAATTGCCAGTTTAGAGGCACAGTGTGG CAAACGTTCCAAGGAATTCAACTATCACCACTTGAATGCAGCGGTGGCGAAAAAT
>Gm03_3699677 7	CATATATTTGCATTCACTTAACGTAGTCTTGATACCCCAATCCACTAACAAATTTTTTTTTAT ATTATATTATTATCATCATCAAAATGCCTGGCTTGTAATGAAATGGATTGGACCAAA CTATCTAGTGACGTGCGCAATGAAATCATCCAGTTCCGCGCGAGAAACACCC[G/T]TTTCAT CCTTGACCTGCGAATAGCATTTTTTCAGGTTTCATTGCCCTTTGTCTCATCTCATCCTTCT TTTGTGGCATCAATCTCCTCACGGCATTCTCAACATCTGATGATGTCACCAACTCGTCCCT GTGGTCCCAGTCCCTCACACCACCCCAACTCTGAGCACTTCTGTTACCAA
>Gm03_3699718 4	GCACCCCCATTGTCATGCTCTCCATGCAAGAGTTCATCCACAGTGACTCATAAACCCACC AGTTGAACTGTGACTTAGAATCTCCAATTGGGGTGCCAGTCTCTCACAACCAACCCCGTG CCTTTCATCTCTCTTCAAACCCCTTTGGAAGCTCAGCAGTTCTCACGCCATCCTCA[C/A]G GAACACATCTCCCTTGTGAGCATCCCTCACCACCCATATGAACTTTTGCTTGTCTTTTTCCA ACCCGTTTGCAACCTCCTTGATTTGCTCCTCAGAGAAGCATGTTGTTGTCCCAAACGACAC ATACAAAACCTGACCCTGCTTCTTGCTTGTCAAGCCACTCAACACTGAAGTGCCTTG
>Gm03_3699718 5	CACCCCATTTGTCATGCTCTCCATGCAAGAGTTCATCCACAGTGACTCATAAACCCACCA GTTGAACTGTGACTTAGAATCTCCAATTGGGGTGCCAGTCTCTCACAACCAACCCCGTGC CTTTCATCTCTCTTCAAACCCCTTTGGAAGCTCAGCAGTTCTCACGCCATCCTCAT[T/C]GA ACACATCTCCCTTGTGAGCATCCCTCACCACCCATATGAACTTTTGCTTGTCTTTTTTCCAAC CCGTTTGCAACCTCCTTGATTTGCTCCTCAGAGAAGCATGTTGTTGTCCCAAACGACACAT ACAAAACCTGACCCTGCTTCTTGCTTGTCAAGCCACTCAACACTGAAGTGCCTTG
>Gm03_3702487 7	TTTCTCATTGTTTTAAAAGGGTGTTCATTGTCCCTTTTCCATGAATTGCAGTTTATGAGCC ACAAGTGCAAATGGGAATGAAATGAAGAATGGGGTTAGAGTTAGTCAAGACAAGTGGTGT GGTGTCTATGTTATGACTGCACACATGTGAAGTGAAGTAGAGACTATTAGTCC[C/T]CAG CAGCTGTTTCTAGTGTGTGTGTCATTGCATTCTCATCCTTTTCTCTTTTTTACGCCTTAA TTTCTTTCTCTTTCTCCCTCTTCTCTCTGGAATTTGGAGCATCAGCCAGCACTCTATGG ATTCTGTATTGGTAATTGGCCATCTACGATCCTCACAACTTCAGTCAGCT
>Gm03_3702495 8	GAAATGAAGAATGGGGTTTAGAGTTAGTCAAGACAAGTGGTGTGGTCTATGTTATGAC TGCACACATGTGAAGTGAAGTAGAGACTATTAGTCCACAGCAGCTGTTTCTAGTGTGTGT GTCATTGCATTCTCATCCTTTTCTCTTTTTTACGCCTTAATTTCTTTCTCTCTT[T/C]CTC CCTCTTCTCTCTGGAATTTGGAGCATCAGCCAGCACTCTATGGATTCTCTGATTGGTAATT GGCCATCCTACGATCCTCACAACTTCAGTCAGCTTCGACCTTCGATCCTTCTAGTTCTTCT GTAAGTTGCTGTTGTTGTTGTCTATGAAATTGATAATCCTGGTAATAATTACT
>Gm03_3702719	GTTAAGCAACCTTTTTGTAGAAGGGTCTTCTATAATAACTCTAGTTCAATTTCTCCAATGTT ATTTTTCAAGTTGCTTAACTTTTGTTTTTTAATTCGGTAGACTCAATGGTGTCTTCTAATTC

8	AAAGTTTGCTAACCTGTTGAAGTTCCAACAGTGATAAGTACTGAAGCCAAAA[G/C]CATCCT CATGAGACATATTTATCAGCATTCTGAGCAGAAGGTTAGTGACTTGATGGTTAAAGAGCAT GTGTTTTGGTGCAGTTAGGTATGCATATGCTTGATGCTCATAACCTTTTGTGTTTTTCAGTT GAATCCAAAAAGAGCTGCATCTGATAACCTTCTTTTACCAGAGCATGGATG
>Gm03_ 3714439 8	TCCTTAAAAAGTTGACACCAAAACTGAAAAAGCATTATCTATGTTGCAAAATGTTTAGGGCT TTCGTAATAATTGCAACAAGCAGATGGGAGTTTTTCAATGACAGGTTCAAGAAAGGTGAAAAG AGAAGTTCTACATGAGATTCGGCGAAGAAAAGCTTGGACCAGCAAGCAACAACCAA[G/A]G GCACCAAACCAAGCAACACTCCAAGACTCTGATGAAGATCAAAGGTCCTCATCAATCTCAT CATCTTCTGGTTACACTACCCTTGTGGATGAAAACAAGCGACTCAAGAAGGAGAATGGGGT GTTGAAGTCTGAGCTCACAAGTATGAAAAGGAAGTGCAAGGAAGTCTTGATTTGGT
>Gm03_ 3716540 9	AAGATTCACACGTACCATCTCGAAACAGTTGTTCTGATAACTGTTCCCTGAGAAGCTCCATCA CAGACAGGTGTACCAGAGATTAGTGTCTTACTCTTCTTTCATTGTCAAGCCAGCGCCCAAT TAAGCCTATCATTTCTCGTGTCAAGCTAATCTCCGCAATGGATGTATTTCTGAT[C/T]TGAC GGAGTCAGTGCCCCAATGCAAATTTCAAGGTCATTAAGTAAGCTTTAGGATGAGAAGGG ACATCAAGAAGTCTTTCTTGAATAGCTTGGCAGAGGTTTTGCAAGTCTGCAGAACTCACTTC AGTAGGTTTGACCTGGAACAGGGAACAGAGAATTACATATTTGAAGATGCT
>Gm03_ 3730000 9	GCTCTCATCGATGCCCAAACAACCCTAGTTGCGAATTTAGAAAGGAAGTCTCGCTGGCG CTCACCATTTCCTCGCGTGCTGCCACGTGGCGTCAAGGCGATGAGGACGAGGCCAGA GGAGGGGCGGAGGGAGGAGATGTGGAGGGCGGAGGCGGAGGAGGGGAAGAGGTAGAG GG[C/G]GGGGGAGAGCGGTGAGGAGGGGGGAGAGGCCGCGGCGGAGGCGGCGGGC GGTGAGGGAGGAGGCGCGGAGGAGGGACTTGGTGAGGATGGGGGTGGTGAGAGCTTG TGGCGGGCCTCGTGGGGGTGTTGGATGATGAGGATTTGGGTGGTGGTTGGATTGGAGG AAGTGAAGCGC
>Gm03_ 3773076 8	GCAGGTGTGTGCACCTTGTGTGCTGAGCTTCCATGCTCTGGTGCAATGGAATAAAGTGACA CTTCAAGAGCAGCAACTTCTCTAAGCTCCTCTTCGAGCTTTTCAATTCTGAATTCCATTCTT CAATTTTTAATTCTGCAGCTGTATCATCTCATGTTACCATTCTTTCAGCATT[C/T]ACTGT TTCATGAATTTTCATCTTCACTTCTTACACTCTGAGCACCATTGGAGGATCCATCCACAACAT CAGGTTCTTGACCTCTTGAACAGGTTTTTCATCTACACCTTCAGAAGATTCTGATGATACA TTTACAGGAACTTTCACAGTCTTGAATTGGTTGAGGGAGTATTTTTGTT
>Gm03_ 3773169 4	CCTTGAAGCTTTTCTCTTCTCAATTTCTTTCATGCTTTCAAATTCCTTCAAATATCAATCAAAT AAAGATGATGTGATAGGGGGAAATGATAGCAGTAACCTTAGCGAGCTCCTTAAGCCT TAGTTCTCAAATCCTAAAGTCTGTTTGTGTAATCACTAACAATACCTACTCC[G/T]TAGAAT AGTGTGTTGGCTTCCACCACCAGAGAAGTGCTTGTAACCTCTCTATCTTTCCAGAGGCT CAAAACAACAATTGTCTACCTGCAAAAGTATTGGTAAATAAGTGGCTTGATTCTCAGAGAA TAATAGACAAAAATCGTATAAGGAAACCAATTTGATTATCGGGATAAGAG
>Gm03_ 3778700 6	TATCCCTGAAGTTTATGCTAAAAAGCAAGTCATTGTTGCATTCCATATTGCTCTAAACTGCA CTGAACCTTGATCCAGAGTTGCGACCTAGAATGAAAACCGTGTGACAGAAACCTTGATCACAT CAAAATACAGTGAGAAACAAGAGATGTCTTAGTGATGAGGGTTGATTTTGTAACT[T/G]GC GGGCTAGTAGATGTATCTCTTCTTCCATTCTCAATGGAAATTGTTTCTTTTCACTTTTATT TAAATCAAGGAACGGCCCAAATTGATTGGTGTGGTAAATATTTTCTTCTTTTACATCCT GTTTACTTCAAATACAAATCCTACGGATTGCGATTGAACACCTTGCTTAAAT
>Gm03_ 3782392 2	GTCTAATTAGTATTATTTCTCTTCTTAGTTTCTTCTATATATGCTATCACATAATCAACAAA AAAAAACACAAACACTTGATAGTACTACTCAAGGTATGGCTTTTCAAGTGTTGTTCAATTTG TTTGATCTCAACCATCGTGTGTCATCCATACTATGGAGAAAAACAAAACAAA[G/A]CAAGAC CCTTCTTCCACCGAGTCCAATGCCCCTACCCATCATTGGACACCTCCACCTTCTTCTCCAA CACCTCACCAAGATTTTCAAGCTCTCACTCCGCTATGGACCCATAATACACCTTTTCTT GGTTCAGTCCCCTGTGTGGTGGCTTCCACAGCAGAAGCCGCCAAAGAGTT
>Gm03_ 3782557 9	GAGTTTAGGCCAGAGAGGTTTGTGAAAATGGGAAGAGTCAATTGGATGTTAGGGGACAA CATTATCATCTACTTCCGTTTCGGTAGTGGAAGAAGAGCATGTCCTGGTACTTCTTTGGCATT GCAAGTTGTGCATGTGAATTTGGCAGTTCTAATTCAGTGTTTCCAATGGAAGGTTGA[A/G]T GTGACAATGGCAAGGTGAACATGGAAGAAAAGGCTGGCATCACTCTTCCGAGGGCTCACC CCATAATTTGTGTCCCTATTGCAAGACTTAACCCATTCCCTGTTGTGTGATTGTTTTTATTT TAAACATCATTTTTCCATTACGTATATGATATGATGTCATGTCAAACATTTATGC
>Gm03_ 3782739 9	GTGAGTGATCCATACCCGCTAGGTTTTACACATTCTTTTCCCTGAAACAAACACCACCCCT TTTAGGTCTTCTTCTAATCCTCCCTCTCTCTAATTTTACACGCGAGCGGCTTCTTCTCCGT TTTCTGAAATCGGTGTGAACAATGGCGACGTACGCTGCGATGAAACCCACCAAGC[T/C]AG GGTTGGAGGAGTCCCAGGAACAGATCCATAAGATAAGGATCACCTTTCTTCCAAGCACGT

	CAAAAACCTCGAGAAGGGTATGCACCTATTTATTCTACTTTTTGTTTTAAGGTTGTTGCTAT CCTTAGCCTTATGTTTCTAGTTTTGTGTTACTGTTAGTTTGTGCGGACTTGGTT
>Gm03_ 3782868 4	CAACACATGGGATAGATTGAACTTCGTGTCCACAAGAGAGTGATTGACCTCTACAGTTCC CCAGATGTGGTTAAGCAGATTACCTCTATCACAATCGAACCTGGTGTGGAAGTTGAGGTGA CCATTGCAGATGCTTGATCTATATCCAAATTTAACTGTTATGGTTTCCTAGTTTTT[G/A]GTC TCCTCTTATTCTGTATTTTATAAGTTTTGATTGCGGGACCGAGTTGAATTGTTTTCTTTGTT TTACGGTTGTAGACGGTTGATGAATACCTGAGAAATGTCATTTTTATATATTCTAGAATCCTA CATATATTTGTTTTGAAATTATTACGAATCATTTCTAATGGACTCAGCTAA
>Gm03_ 3783222 8	CCTTGTGTTTCATGAACTGCCTTTTTGTTTTAGGTACCAACACATGGGACAGATTTGAACTT CGTGTGCACAAGAGGGTGATTGACCTCTACAGTTCCCCAGATGTGGTTAAGCAGATTACCT CTATCACGATTGAACCTGGTGTGGAGGTTGAGGTGACCATTGCAGATGCTTGATCT[C/T]GA TCCAAATTTAACTGTTATGCTTCCTAGTTTTTTTCCCTCTCTTTTGTATTTTGAAGTTTT GATTGTGGGACTGAGTTGGATTGTTTTCTAGTTTTACGGTTGTAGACATTGATGAATACC CGAAGAAATGTCATCTTAATATATTCTAGAGTGTTACATATTTTGTGTTGAAAT
>Gm03_ 3786380 4	GTCACCTTTCTGTCTCCAGCCTATACCACTGCGGTGCCAAGGGACTGTCTGGTGGAAC ACGCTTTGGGATCTCATTGAGGTCAAATAAGACACGACCAATGAAGTCATCTTTCACGACA TCCTTGTCTTCACAGTGACCTCCAGTATGGAAGCCTGAATGCGGTCTTTGGAGAAAG[T/A] AAAAACCTGGTTCCATTGCGGATTAGACTTCTTGTCGAAGTGCCGAGTGGTGCCTTTGTAG TTCCCCAGCTTGACTTCAGTATAAGGGTCACAACCTGCCAGTGACATCCTTTCAGGTAAGT CCTTAGCTTTCACAACCCTCACATAGAGATACTGCATTTGCTCGACTAGGTCATAAGT
>Gm03_ 3805524 0	GATATATATTGACTTCCATGTTTTATTTTCTTATTAGTTTTCAGTGTAACATAGTATTTATACA TTAGTTAAAAATTGGTTTGCACAAGGATGGAAGTCAGTATTATTGGTTTGGTCTCAACCTGG CTGGTTCAAGAGGAATCAATCATCTGATGGTAGAATCTCACTCTTTTAGTCAT[A/C]AAGATG ATAAAGGATGGCTGCTTAGACACCCACCTGAGTCCGCCCGACTAGTCAAATCTGTTCTTTC TTGGAATCACGTATTATTTCAGAGAGGCCAACCAAGTAGCTGATGCTTTATCCAAACATGGT CTTTCCTTGGTACTACTGATAGCATTGTTTTTTTTTTTATTACAGTTCCGTCT
>Gm03_ 3805529 0	ATAGTATTTATACATTAGTTAAAAATTGGTTTGCACAAGGATGGAAGTCAGTATTATTGGTTT GGTCTCAACCTGGCTGGTTCAAGAGGAATCAATCATCTGATGGTAGAATCTCACTCTTTTA GTCATTAAGATGATAAAGGATGGCTGCTTAGACACCCACCTGAGTCCGCCCGACT[G/T]GT CAAATCTGTTCTTTCTTGGAAATCACGTATTATTTCAGAGAGGCCAACCAAGTAGCTGATGCTT TATCCAAACATGGTCTTTCCTTGGTACTACTGATAGCATTGTTTTTTTTTTTATTACAGTTCCG TCTTTTCTTCTGTTAAGATATGTGCTGATGTATATTGTCTACGACCTTCCCT
>Gm03_ 3806480 2	TTCAAGTTTCTACTTGCTATTGAAATACAAATTACCAAGTTATTTAATGTACTGTATTAACATT TCAAAATTCAACAGCTTCAGGCATATGTAATTTATAGTCATGCAGTTTACCTGTTGGAAGGT GCTGAGGGAGGCCAAGATAGAGCCTCCAATCCAGACACTGACTTCCTCTCTG[G/T]TGGT GCTACAACCTTAATTTTCATGCTACCCGGTGCCAAATGCTGTAATCTCCTTGCTCATCCTATC AGCAATACCCGGGAACATTGTGGAACCAACCACTCAAGACTATGTTACCATAGAGATCCTTC CTGATGTGACATCACACTTCATGATAGAGTTATATGTTGTCTCATGGATACC
>Gm03_ 3806506 6	TGTGGAACCACCACTCAAGACTATGTTACCATAGAGATCCTTCCTGATGTGACATCACAC TTCATGATAGAGTTATATGTTGTCTCATGGATACCAGGAGATTCCATCCCAATCATGGATGG CTGGAAGAGAACTTCAGGGCACCGGAATCGTTCAGCACCGATCGTAATCACCTGCC[A/G]A TCAGGTAGCTCATAGCTCTTCTCAACTGCTGAGCTGGTCTTGGCAGTTTCCAACCTCCTGCT CATAATCAAGAGCAATGTAGGCCAGTTTCTCCTTCATGTCCCTCACAATTTCCCGCTCCGC AGATGTGGTGAAAGAGTAACCACGCTCAGTCAAGATTTTCATCAGGGCATCAGTGAG
>Gm03_ 3808341 7	GTTTTCTTTTCTCTCCATCCATGGATTCTCTGAGTTCCATCAACGGCGATTTCGGCTTCGA CGGCGGCACGGATCGCAGATTTCGCGTTTTTCGCGCCAAGCCTCCTTCCAACAACCTCACAC GCCGATAGACATTCCGCGCACGGCCACTACGACCACCACCACTACTGGTCCCCGCG[A/G] JGACGATAAACCTCCGCGCCTCTCTCCAAATATCCTCTCTCTCCTCCTTCGTTTTCTCCG TTTTCCGAAGCGTTAGGTCCGCGCCACAGGTACATGAAGAGGCTCTTCTCATGATCTCGCT CAATGTGCGTACTCCACCGCCGAATTGCTCACTGGACTCTTCAACGGTCGCGTAGGT
>Gm03_ 3811745 3	GTACATGAAATTATAATAACCGTTTTCTGCACCTGTACTTAATTATTGGCCTTACTTGCTATT CTCATGAGCTACAGCAGCAATGTGGTTGCTTGTGTTGCCAGCAAACCTGCTTATGGAACGC GCTTTACCATGCCAAGCTTCATCCTCTTCTCGTCGACGTCATTATTATCCCTG[G/T]CTTC TTCTCCTCCTTCTCCCAAAGGGCTCAGCATCATCAACACACTCAATTTGCTTCTCGAGAGA ACAAAAAGAAGGGGAAGAAGAAGAAGGGTGAGGCACGTGGTGCACGTGCACCTCCTTTGG TGCAGTTGATGCAATAATGGAACGGCTTCTAACACTTGTATGCTTCTGAGTCT
>Gm03_	TTGTACTTAATTATTGGCCTTACTTGCTATTCTCATGAGCTACAGCAGCAATGTGGTTGCTT

3811748 5	GTTTGCCCCAGCAAACCTGCTTATGGAACGCGCTTTACCATGCCAAGCTTCATCCTCTTCT TCGTGCGACGTCAATTATTATCCCTGGCTTCTTCTCCTCCTTCTCCCCAAAGGGCTCA[T/A]CAT CATCAACACACTCAATTTGCTTCTCGAGAGAACAAAAAGAAGGGGAAGAAGAAGGGT GAGGCACGTGGTGCACGTGCACCTCCTTTGGTGCAGTTGATGCAATAATGGAACGGCTTC TAACACTTGTATGCTTCTGAGTCTTCCCTTCCCCAAGTGATGCTCACACAGCGAGT
>Gm03_ 3811817 5	TCCCATCTACAACGTATGCAACACCAACACAGTAAACAAAAAAGGGGATCAACAAATAAACCA CAGCAAGCAAGACACAGTTAGTTAGTTAGTTGCATTACATGCATTGAGTTTTTGGCATTAC CTTGAGTAGAAGATGAGCTTAACGGTGGCAACAACCTCAGCTTGTGTTTTTCCGA[A/G]TCA AAGGTGCGTCCCTTTCCTACAAACAAACCAGACCCAGAAAGACCCACAATTCATAACATAC ATTTGCACATACTTGTACGTGAATACATAAAATTTACGGCAAAAGGGTGTGAATTAAGTGA AAGTTGGAGTGTAATATTTGGAAGGAACAAAAACGCAGTGTAGTTGGGGACCAG
>Gm03_ 3811958 4	AGGTTTCAGATTTCAAAACAAAAATAAGATAAAGAAGAAAAAGGGTATCAACAAATAAACCA AAAATCTATAGCTACTACATCTACATGACCCTCCAAAGACCAAGAAGAAAAAATAATTATGT TTTTATCAACAACCTTTTCTCTCTCTATCTCTCTATCTCTCACACAAACGCAT[G/A]AATAC ATACACTCTTTTCTTCTCTCTCTCTCTATCTCTATAGCACTCCATATATATTGCTTCCAT GAGTTATTATTCTCTCTCTAACTTTTATTTGTTGCACCACAGAAAGGGAGTTGTATGGGTGA GGACGGCACGTGCGATGGAGTTAAGGGAGGAGGAGGTACGTGTGGAACATA
>Gm03_ 3813299 6	TGACAATTTACCTATAGAAAAATGGTTGATTTAATTACATAATTACTATGATGCTTATTAAGT AGTGGTTTTGTTTTGAGTGGAAAAGGTGCACATAGTTTTGGTCTGCGCAGTGATAGTTTTT CAACCAGAGACTGTTCAACTTCAGCGGCACAGGTAGCCCTGATCCACGCTGAA[C/T]ACA ACCTATTTAGCGACACTTCAGCAAACTGTCCACAAAACGGAAGCGGCAACACTCTGAACA ACCTCGACCCCTTCGTCTCCGGACACCTTCGACAACAACCTACTTCCAAAACCTTCTCAGCAA TCAGGGCCTTCTCCAAACAGACCAAGAGCTCTTTTCCACCAACGGCGCCGCCACA
>Gm03_ 3813335 9	TCCGTCGTCAACAACCTTCGCCGCCAACCAAAACCGCGTTCTTCCAAGCCTTTGCTCAGTCCA TGATTAACATGGGTAAATATTAGCCGTTGACTGGAAAGCCAGGGTGAAATCAGATCAGATTG TAAAAGGGTCAATGGAAGTTAAAAATGAGAGGCACAAGCTAGCTGTTCTTGATACAG[C/T]A ACTCAGTTACGTTACGTTCCATTCAACATTTACTTAAAACCTTTGCGTTGTTATTTGTGTGATC ATGTCAAGAACTCAAGACATGGTAGGATGTTTAAAACGTTATTGATTTGTATGTTTGACTAT ATATATATATATATATATATATATATATATATATATATATATATATATATATATATATATAT
>Gm03_ 3815018 1	ATATATATACAGTGGGTATGGAACGTGTGACATCAGAGATTCCGGTGAGGGTGGTGGATG ACACCGACGATGGTGCTGCACCTTTCATCTCCCAACAGTGCAGTGTCTCTTTTCAGATGGA TTATTGTGTCAGGAACAATAGAAAGTCTAGTGAAGGTGCCAGCGATGACGACGAAAAT[T/C] GCTCATCAAGGAAGAACTAAGACTCTCCAAACAACAATCAGCTTTCCTCGAAGACAGCTT CAAAGAACACACCACTCTCAATCCCGTAACCTTTCACCTTCTCTTTCTCTTCTATTATTTT TTTTAATCATTGAAAGATAGAATAATATCTATAATTTTATTCAAGTTTAATTATA
>Gm03_ 3817043 1	TTAGATATCTTAGTATGGGAAAAATAAAAGAAATGTGAAATCTTTGCAGCTGTGGAAAGAT GGAGAGATGAAAGAAGAAGTCATTGGAGGCCATAAGGCCTGGCTAGTTATTGAGGAAGTC AAAGAAATGATCCAAAAGTATTTATGATTCTTCTATTTTATTTTTCTTGCTCCCTCA[A/C]CCC CCCACCCACCTCACAAAAAAGAAAGAAAGAAAGAAAGTTACAATTGAAGAACTA ATCTTCCAATGTTTTTACTTTTTTTCTTTAACATTTTTCATGTTGGCAACCGCATGTAGCTGT GTATCATATATATATATTAATGGAACGTGTGTACCATAGCTGAAAGAAATTTA
>Gm03_ 3817052 4	ATAAGGCCTGGCTAGTTATTGAGGAAGTCAAAGAAATGATCCAAAAGTATTTATGATTCTTC TATTTTATTTTTCTTGCTCCCTCAACCCCCACCCACCTCACAAAAAAGAAAGAA AGAAAGAAGTTACAATTGAAGAACTAATCTTCCAATGTTTTTACTTTTTTTCT[A/G]TAACA TTTTTCATGTTGGCAACCGCATGTAGCTGTGTATCATATATATATTAATGGAACGTGTGTAC CATAGCTGAAAGAAATTTAAGTGAAAAAGGAAGATGCTCACTGTTTCTCCCTCCCTCTCAA CCATTAGAGATTTAGAGGGGAAAAAAGTGGGGGTTTTTATACCAATAA
>Gm03_ 3817352 5	ACCTCCTTACTAATCCTCGCATTGCTGTGTAAAACCTTCCACCTATTTTCATCCATCAA AATCTCAAACAAATCTTAGCTCTGCAATTAACCTCTCCGTCATGCTAATACTCTTCTTCC CTTCTTACCTTCTCCTTCTTCTCCTTCTTTCGGCGCAGCAGCCGTTGTTTT[A/T]GTCGT CGTCGGAGTCGTGTTGTCGTTGTCGCCGCCGAGACGGCGACAACGACGACGGCGCAA CCTTCTTGGGTTCCAATTGCTCCTTAACAGGACCTCCTTTGGCCATGCTCTGCACCCAAAC CCTAACCTTCTCCAAGATCAAGGGCTTCCCCTTGGAACCATGGCATCCTTAATC
>Gm03_ 3817371 9	CGTCGTTGTCGTTGTCGCCGCCGAGACGGCGACAACGACGACGGCGCAACCTTCTTGG GTTCCAATTCGTCCTTAACAGGACCTCCTTTGGCCATGCTCTGCACCCAAACCTTAACCTT CTCCAAGATCAAGGGCTTCCCCTTGGAACCATGGCATCCTTAATCCTCTTTCCAACCG[G/ A]TCCCCTCGTCGTCGACAGTAACCCTAACCTCGGGATCCTCGTCGGCGTTCTCGTCGGAGA

	TGTAGGGGATCTCGACGGTGCCGTCGACCTTGAGAAGAGAGGTTCCCAGGGTGTCTTTGG CCTCTCCCTGCCAATTGAGAGTGAGGCTGATCTCGTAGCCGGGGATGATCTTCCCCTTTCCG
>Gm03_ 3817381 5	GCTCTGCACCCAAACCCTAACCTTCTCCAAGATCAAGGGCTTCCCCTTGAAACCATGGCA TCCTTAATCCTCTTTCCAACCGGTCCCTCGTCGTCGACAGTAACCCTAACCTCGGGATCCT CGTCGGCGTTCTCGTCGGAGATGTAGGGGATCTCGACGGTGCCGTCGACCTTGAGAA[C/T] AGAGGTTCCCAGGGTGTCTTTGGCCTCTCCCTGCCAATTGAGAGTGAGGCTGATCTCGTA GCCGGGGATGATCTTCCCCTTTCCGACATTGATGTAGGCCTCGCCGTCGAGGGAGCGAAG CGACGTCGTTTTGAGGAAGAGGTTACCCTCGCCGTCGAGGATAGGGAGGTTAGTGAGGAG
>Gm03_ 3817415 0	CGAGGATAGGGAGGTTAGTGAGGAGGCTGGTGAAGAAGGTCTTGACCAGTCGAGGCAGT TGGTCTCGGACCAATGCCAGTTGTGGACATTGGTGCCGTCGGGGCGGTCCCTCCACGATCC ACCGCTTGTGCGCCCTCACCCTAACGAGCCATTGCAACTGATGCCTCTTTTCTGGGAGAGT[ T/C]CCTTAGTGATTAATAATAATTGGATTGGGGTGAGGGTTATGAACTATGATGATTATTATT ATATGTGTTGAGTGTGTGTGGAATGGTCTAGAATGAAAAGGAAGGGTCTAGAATGTTGATG TTGCTTTGGGTACACTTTTCTCACCAACCAACCAAGCCCGAGAAGGTTCCCTAGGAGTT
>Gm03_ 3817415 7	AGGGAGGTTAGTGAGGAGGCTGGTGAAGAAGGTCTTGACCAGTCGAGGCAGTTGGTCTC GGACCAATGCCAGTTGTGGACATTGGTGCCGTCGGGGCGGTCCCTCCACGATCCACCGCTT GTCGCCCTCACCCTAACGAGCCATTGCAACTGATGCCTCTTTTCTGGGAGAGTTCTTAG[ G/A]GATTAATAATAATTGGATTGGGGTGAGGGTTATGAACTATGATGATTATTATTATATGT GTTGAGTGTGTGTGGAATGGTCTAGAATGAAAAGGAAGGGTCTAGAATGTTGATGTTGCTT TGGGTACACTTTTCTCACCAACCAACCAAGCCCGAGAAGGTTCCCTAGGGTTTTCTTTT
>Gm03_ 3838944 9	TATGCCCAAACACACAAATAAAAAAGAGAATAGATATCTCGATTACAGAGAACTCAAATAAA TGAAAACTATGCAGGAACCATGTTAGAAAACAATAATTATTTTCAAATGTTAAAGGACTAT ACATCATCCAATGTCAATCCTAATCTTACTATATAAGGCAACCAACTACATACGA[T/C]AACA AGAGCCTTAAATCTAAAACTCCTGACTAGCAGATCCAATCTCGGTCTTCCCCTTGCCAGCG ACCTTCTTGGGCAACAAATTCTGGTGAATGTTAGGCAGAACACCACCGTTGGCAATGGTCA CAGATCCCATCAGCTTGCTCAACTCCTCGTCGTTCTTAACCGCCAGTTGAATGT
>Gm03_ 3838945 7	AACACACAAATAAAAAAGAGAATAGATATCTCGATTACAGAGAACTCAAATAAATGAAACT ATGCAGGAACCATGTTAGAAAACAATAATTATTTTCAAATGTTAAAGGACTATACATCATC CAATGTCAATCCTAATCTTACTATATAAGGCAACCAACTACATACGAAAACAAGA[A/C]CCTT AAATCTAAAACTCCTGACTAGCAGATCCAATCTCGGTCTTCCCCTTGCCAGCGACCTTCTT GGGCAACAAATTCTGGTGAATGTTAGGCAGAACACCACCGTTGGCAATGGTCACAGATCC CATCAGCTTGCTCAACTCCTCGTCGTTCTTAACCGCCAGTTGAATGTGCCTCGGA
>Gm03_ 3871767 3	AAAAATAGAAACAATCAAATTACCTATACATATTTTTGAATTGGAATTTATTTAATTCCATTT GCAAAATTATTGACCTTGAGATTAGAACATTTGGCATCCTGAAGCAAGAGAGTCGAAGTGA AACCGAATCTATTTGAGTCAAGATCAAAGTGCATTAAATTATCCTCCAAGTAT[C/T]CCCTC CGATTACAATGGGAGTTCTCCCGCGGGTCCACCGTCCACAAAACCCAAACACCACACAT CCACTCCTCCCTTCGCAACCCCTCAGGATCGAATTCCTCGCAAAAGATTCTCCAGAACACGTC CTCGCTGTGCATCACCAGATCAACGGTGGGCACCGCGGGTCCAACCTCGTGTCTC
>Gm03_ 3871768 2	AACAATCAAATTACCTATACATATTTTTGAATTGGAATTTATTTAATTCCATTTGCAAAATTA TTGACCTTGAGATTAGAACATTTGGCATCCTGAAGCAAGAGAGTCGAAGTGAACCGAATC TATTTGAGTCAAGATCAAAGTGCATTAAATTATCCTCCAAGTATGCCCTCCGA[G/A]TACAA TGGGAGTTCTCCCGCGGGTCCCACCGTCCACAAAACCCAAACACCACACATCCACTCCTC CCTTCGCAACCCCTCACCATCGAATTCCTCGCCAAAGATTCTCCAGAACACGTCCTCGCTGTG CATCACCAGATCAACGGTGGGCACCGCGGGTCCAACCTCGTGTCTCCGTGAGATC
>Gm03_ 3871867 2	GACGGGGTTTTCGGGGAAGAGGGGCGCAGAGGGAGGAGTTGTTGGAGCATGCGTTGGAGG GGAAGGAGTTGCAGAGAGGGGTGTTGCAGGGGATGTGGTGGGAGGAGGAGGAGGTGTA GGTGGAGTCGCAGAGGACCCAGCTGAGGGAGGAGCCAAGGTGGAGGTGGAGTTTGTGG GTT[A/C]GAGTGGGGTTTTCAGGAACACGGAGAGAGTGTAGAGTTGGGTGGTGTCTTTT TGAGATTGGGGCTATGAGGGGGATTTGGAAGTGGGTGTGAGGAAGAGGAAGAGTATCGC CAAGTTGAAGTTGCAGAGAGATGGAAGAGGAGGAGGCATTTTGGTTTGCTTGCTGGAGCA AGGAGA
>Gm03_ 3871871 0	TGTTGGAGCATGCGTTGGAGGGGAAGGAGTTGCAGAGAGGGGTGTTGCAGGGGATGTGG TGGGAGGAGGAGGAGGTGTAGGTGGAGTCGCAGAGGACCCAGCTGAGGGAGGAGCCAA GGTGGAGGTGGAGTTTTGTGGGTGGAGTGGGGTTTTTTCAGGAACACGGAGAGAGTGTAGA GT[C/T]GGGTGGTGTCTCTTTTGTAGATTGGGGCTATGAGGGGGATTTGGAAGTGGGTGT GAGGAAGAGGAAGAGTATCGCCAAGTTGAAGTTGCAGAGAGATGGAAGAGGAGGAGGCA TTTTGGTTTGCTTGCTGGAGCAAGGAGATAAACAAAGAAACAGTGGTTGGTGTGGAGTTA



	AAGTG
>Gm03_38719968	TTGAGCTTTTTCCAATATGATCTTGTGAACCTTGATATCTGCAGTTAAAAAGAAATACCAAAA AGGAATGAAAATAAAAGAAATAATAGGCAAAATTGCTAATATACAACAGCGACCAAAATCTT GTTAGCCAAACAAAACCATATAACCTAAGCTTCTTTCCTATGTGATGCATGAGC[C/C]GAAC AATGCCGTAGCACAAATACAATCTCCTAGAAAAAATTGTACAGCTTTTCTGCTGGCACTGCA TTCAGGAGTCTTTGCTAGCTCTTCAATAAAAAATACGATGTCGTTTCGACATGCTTAGCTTTTA TTATAACTCACTTGCGCCGCGAAGAGATCATATGCTGGTGAGCCTTGATATC
>Gm03_38720115	ACCTAAGCTTCTTTCCTATGTGATGCATGAGCCGAACAATGCCGTAGCACAAATACAATCTC CTAGAAAAAATTGTACAGCTTTTCTGCTGGCACTGCATTCAGGAGTCTTTGCTAGCTCTTCA ATAAAAAATACGATGTCGTTTCGACATGCTTAGCTTTTATTATAACTCACTTGCGCC[T/T]CGAA GAGATCATATGCTGGTGAGCCTTGATATCCCTTTGTATTCTGTCTGCATACAAGGGTAGGC GAATGATCTGATTGATCTCATTCAATAAGACCTCTTCTGTGATGTTATCCTTTATGCTTCGCA TTACCTGTGTTAAAAAATTAACAAAACTGTTTCAAGAGGCTTTTACATGAG
>Gm03_38942898	ACGGAGTTCAATTTACTGAATCTAATTTCTTTTTTTTTTCGCATTTCTATTTTTCTCCTCATT TCCTTGGGAACCAACAGTTAGATTAATGACAGTTAGATTGATGAGGTACCTGAATCCTGA CATAGTCGGTACGATTCCAACAGAAGGCGTTCCCTAGCATCTGGTCAACGGTCT[G/G]TGA AACACCGTCGAGCGCAATGTCAATCAGCATTGACGTAGAGCACTCGCCGGCATTGTTTCATT CTCTTCGCCGGAGCTCCGTTTCCGATGGAGAGAACAAGGAGATCCTCCACGCCGTTCCACC GACGGGAAGTCGCGCTTGTGTGGAGGACGTGCGTGACCGCCGCCGCGGCCGGATT
>Gm03_38942920	TAATTTCTTTTTTTTTTCGCATTTCTATTTTTCTCCTCATTTCCTTGGGAACCAACAGTTAG ATTAATGACAGTTAGATTGATGAGGTACCTGAATCCTGACATAGTCGGTACGATTCCAACA GAAGGCGTTCCCTAGCATCTGGTCAACGGTCTCTGAAACACCGTCGAGCGCAAT[G/G]TCA ATCAGCATTGACGTAGAGCACTCGCCGGCATTGTTTCATTCTCTTCGCCGGAGCTCCGTTTC CGATGGAGAGAACAAGGAGATCCTCCACGCCGTTACCGACGGGAAGTCGCGCTTGTGTG GGAGGACGTGCGTGACCGCCGCCGCGGCCGGATTGTTTCATCACTAAGCCGCCGCTCG
>Gm03_38942927	TTTTTTTTTTCGCATTTCTATTTTTCTCCTCATTTCCTTGGGAACCAACAGTTAGATTAATG ACAGTTAGATTGATGAGGTACCTGAATCCTGACATAGTCGGTACGATTCCAACAGAAGGCG TTCCCTAGCATCTGGTCAACGGTCTCTGAAACACCGTCGAGCGCAATGTCAAT[G/G]GCA TTGACGTAGAGCACTCGCCGGCATTGTTTCATTCTCTTCGCCGGAGCTCCGTTTCCGATGGA GAGAACAAGGAGATCCTCCACGCCGTTACCGACGGGAAGTCGCGCTTGTGTGGAGGAC GTGCGTGACCGCCGCCGCGGCCGGATTGTTTCATCACTAAGCCGCCGTCGACGGCGG
>Gm03_39790817	CGTGTTCGCCATGCAGGGTTTTGGAATTTTGGCAGGTGGTATCTTTCGAATTATAATTTTCAG TTGCATTCAAGGAAAGGTTTGATGCTCCACCATATGAGCTTGATCCAGCTGGCTCAACTGT TGCACAAGCAGACTACATTTGGAGGATAATTGTTATGGTGGGAGCACTGCCAGCTG[C/C]G TTAACTTACTACTGGAGGATGAAGATGCCGGAAACTGCCCGTTACACCGCTCTAGTCGCCA AGAACACGAAGCAGGCTGCAGCAGATATGTCTAAGGTTCTGCAGGTTGAGATTCAAGCTG AACCGCAGAAAGAGGAGCAGAAGGCTAACTCATATGGCTTATTTTCAAAGGAGTTCTCT
>Gm03_39790874	TTGAGTTGCATTCAAGGAAAGGTTTGATGCTCCACCATATGAGCTTGATCCAGCTGGCTGCTCA ACTTTGGACAAGCAGACTACATTTGGAGGATAATTGTTATGGTGGGAGCACTGCCAGCTG CGTTAACTTACTACTGGAGGATGAAGATGCCGGAAACTGCCCGTTACACCGCTCTAG[T/T]C GCCAAGAACACGAAGCAGGCTGCAGCAGATATGTCTAAGGTTCTGCAGGTTGAGATTCAA GCTGAACCGCAGAAAGAGGAGCAGAAGGCTAACTCATATGGCTTATTTTCAAAGGAGTTCC TCCGTCGCCATGGACTGCATCTACTTGGTACAGCAAGCACATGGTTCTTGCTTGATAT
>Gm03_39963852	AGAAATAAGCTAAACATGCATCAAATTAACATATAACAAGAAAAACAGAGCTAGAGAGGGTT CAATTGACCTTGAAAACGATCCATCGAAGAGCTTAACGCGGCGGAACGGTGAGCATCGG AGACGGGGCCGGGAATCGTGACGGCTCCGCAGATCGAAACAGCGAAAAGTAGCGCCA[T/T] JTAATCCTCCTAGGTTTCGGGCCATTCTGCACCGATCAATGGAAGAATTTCAACAACAAAG CGAATCGCATTTACAATTTGATGCGGATCGGCTTATGTTAGGGTTTGAGATCGAGAAGAA GAGAATTAGATCATACGCGGAAGCAGCAAGAAGCTGAGAAGAAATGCAATTGGCAAGC
>Gm03_39986148	TTGTGAAGATGCTGCATGCAATCATTGTGTTGAAATTTTTATGGCTGGAGTTTATTTTAACTT TCTACTCGTGTGTTCTTAACTTTTTATTTTTATCTTGTCTCTTAAGTATCATATCAAGAGAA GGGACGCAATTCGTCCCGATATGGCAGCAGTTCAAGAAGACCCATAATCTCA[G/G]CGTCA ACCAGGCCAAGTTCCTCTGGTGATCATACTGACAGTCGTAAGTGGCCGGCTAACCTCAAGTG GAAGCCGACAATCTGCCACACATAGAAATATTCAACCTATGCACGAGACAAAACAACCAAC TTACACACGCTCTGGATCCACCAGAGGCAACCGTGATGATCCTCTGCGGAGTT
>Gm03_4002172	GAGTTTCTTGACACTGTGACAATTGTTGGGGGTCCAAAGTCAAGGAATGAAGTGTGGACAG TGTTTGCCACTGCACATCAGATGACAAGTTGTAGAATGGTTGCCACTGAAGAAGAAGGTGC

4	ACTTGATGAGAATGGTGAGAGTTGGGAAGCAAAAGGCTTGTATGTGTGTGATGGCAG[G/A] GTTTTGCCAAGTGCAGTTGGTGTCAACCCCATGGTAACCATTGAGTCTACTTCTTACTGTAT TGCCACTAAGATTGCAGAAATCACTTAGCAAACCATGAAAAATCAAGTTGTACTGTTTCACGA AAAGAAAAAATACTATGCATTCTTTACACTGTATCCCATAGAAATATTATTTAT
>Gm03_ 4005607 0	AAACAATTCTTAATCAACTTCTAGTAGTTTTAAAACTGGCAAGCTTCCTCGGAACTTAACATA ATTTGAATGAGGTTGGGGTTGGCCAACTTTCAAATCCAAGAGTGTGGTGCATGATAAAGC TGAGGAGCTTGAGTTGTCTTGTGGGAGAGGGAAGCCCAATCCAATGCAATCTTCC[C/T]AG ATGGGACTTGGGGGCCGAGAACAGTGTGGAGTTATGGAAATGGATGTGAATCAAGTAACA CAATCAGCACCTGATAGCCCGCCCTTTTGTGATACCAAAGGTTTCAGATAATGACTGCAGTG ATCAAGATAGTGAGCATGTAAGGAAACATACTCTCTTATTTATGGTCTTCTGTTTGT
>Gm03_ 4005979 9	AAAGCTATTCAATGGCTTAAAGTTTTATTTTTCTGGGGATTATGTGTCAACTTATAAGGAAGA TCTCGAAGAATTAATTGAAGTGGGAGGAGGTACTGTTTTGAGAAGCAAGGAAGAATTGGAA TCACAAAGACATGAATGCAAAGGAGATTCTTCACAGTTGCTGATTGTTTACAACC[T/C]TGAT CCTCCTCAAGGATGCAAATTGGGGGAGGAAGTTTCAATTCTTTGGCAAAGGCTAAACGATG CAGAGGATTTAGCTGCTAACACTTTACAAGTCATTGGTCACACATGGATTCTTGAATCAATT GCTGCATGTAAATTGCAGCCTTTTGTAACTAAAATGCATTAGGTAGCTGATA
>Gm03_ 4008404 6	CAACAAAGCATGTTTGATTTTTGGGATTGAAGTTTATTTATGATTCTGATAAAATGGGTTTGG GATCTAGGGCTAAACCTTTGGAGCAGATAAACGTGAGCCCTAAAAAGAAAGTAACTTTTTGA TGTCATGTGAAAACGTACGAGCCCGAGCCTGATGAAGTTGCTGATTACCAACTT[C/T]TGA GAAGTGAAGAGGGTAGAGAGGAAAGTGCTTTTGTGAAAAGTTAAGCCAAACAAAGTCTTA CTCTGAAGTAAGTTCCGTGACTTCAGCGAGGTCTTACCCTACAAACCATAGGTACCACAGT TGCACGTGTAGTGATGATGAAGATGGAGCAATGGAGTATTGGGATAGTGATGTTA
>Gm03_ 4011792 4	GGAACAGAAAGTGACCCAATGCGAATTGGACCCAACAATCAACGAAAAATGAAAAATTGAAG CAAAGGTATTTTCGCAAATTATAGATAAGACAGAATACGACCGAAATTTACGGAACCGGT CGCGTCAAAAATCTTACCCGAAAAGGGGTTACGTGAAGCTCAAGAATTGGATGCAG[G/A] CTAGGGCGGGTGCTTTTAATTCCATCACCGGCAAGGCGAAGAAGGTTGGGTGGGTGCTCG CGAGAGTGAACGAAACGGTGACGAACGAATGGGCTTTGGCTTTGGCTCCTTTTGGTTGGA GGAATAGCAGCTGGCAACCCTACGCGCCAAGGTGGCGCGTGAAGCTCACTATCGGTGATT
>Gm03_ 4013135 0	CAATGAGTAAAGCAGACTATAAAAAAGAAATTAAGATCACAAGAAAAACCAAGAAAAACACCT CTTATAACTATCAATTAATCTCATCAAAACACTTCTCACTGAGAAGTCAAGCAAAAACAACT ATATGTGCCAAATGCCAATATTGTCTGCAATTCAACCTAAAACAATATCTGAAA[A/T]AACTT CAAGAAGCAGAAACATTACATAAACTAAAAACCCAGCTTAAACAATAGCATTTCAGCACTT GCAATCCTGGGCCACAGATCAGCACACTCCTTCCCAATAGGACCAGTGATAGCAGAACCT ACAATTAATAATCTCACCAATCAAACACTAAACATTTACAGAGGAAAAAAA
>Gm03_ 4017373 9	CAGGAATTTAAGAATTTTGGCAGAATTAAGCCTGATGGAATATTCATTAGGGTCCGAAAGG TAGCTCCTCTCTTTGGATTACACTGAATGCTCATATATCCATCTATGCCAAAAAAAAGAGA AAAAAACTAAAGTAGAAATTGATGTGTACAGGAAATTGGAGTCTGCTATGCATTTG[T/G]AG AATTTGAAGACATTGTTGGTGTTCAAAATGCACTTCAGGTTTATAGTTATTCTATTTATTGTT TTGTTTTTTGTGTTAAACCTAATATTTATTGATTATCTAATCAATGATTACCTTAATTAACAG AGAACTAATTGAACATGTCTGAATCATTTATTGCTGATTTGCTTTACTCT
>Gm03_ 4017970 1	CCACCATTGAGCACACTGAACCCATCTTCACTCTCACTTTCCATTTCTTCATCTTCTCTTCT TCACTCAAGCCATTGTGCTTGGCAGTAGAGAAAGAAGGAATGGTCTCCAGCACCAAGCTC CTGTGTCTAGCTGAAAACCTCTTGGAAGGGTCTGAAAACCTCAAACCTTTCTTCATT[T/A]TTG CAAGAAAAAGAAGGCCAAGTCTTGCAAGTTCCCATCAGATTGGACTCTTCTGATCAAGCCTT TGTTTTGCCTTTCAAGCTCACCAATTGAGGGAGAGATTGGAGAAGAACCACATGCAGAGGA CCCAGTTTGGTGGAAGTAAGTTTGTGATAATGTTGTGGGACAGTGGTTGGTGGA
>Gm03_ 4019584 7	GGAGACTGTTTCATTCTTCCGGATCATCATCTTGGCTAAACCGAAATCTGCTATCCTTGCT CAAGATTTGCATCAAGCAATATGTTATTCGACTTGATGTCTCGATGGATAACAGGTGGATGA CAATCATGATGGAGATAAGCAAGTCCTTGTGCAACTCCTAGAGCTATGTTGTACC[C/A]CGA AACCCAGTCTACAAGCAATCTTGTGCTTGTCTACCATGCAAGGCTTCTCCAGGTTGCCA TTGTGCATGAATTCATAAACTATCATCACATCAATGTCATTGTGAATAAATCCTAATAGTCTA ACAATGTTCTATGCCTTAGCCTCCCTAAAACATTACCTCTCCAACAAGGTC
>Gm03_ 4033853 9	CATGCTTACTTCATACTAAAGATTATACAGAATACAAAGAGTGGTTAGATGTGAGAGCAATT CACATCAGAAAAATTTAAGATTGAGAGAGAGAGAGAGCATTCATTTTACCTCTAACACTGTA TATATAGGCAATGTTGTTTCTCCATCGATAACAATACTTTTAAGAAGAGAACCAT[A/C]ACGC CGACCATACGCAAGTAGTATGTGCTCAGATGTAGGCGAGAAGTATATCAAACAGCAATTCA ATCCTCTAAGAGAAGGCCATTTCAAATTATATTTGACCAATTTAATAAACTTGGCCCTAA

	AAGTTGCAACATGAGTTACTTTGGTTGCTGAAAGTGCATGTCTATACAAATCA
>Gm03_40340465	TGTTGGCTTCCTTCAGCAGGTCTGGTTTCATCAGTCTCCATACCATTGTGAGATTACTGAGGCCAATTCCCATTCCATTAGAGAAAGCAGTATGAGCAGGATATTCTGCACCAGTTTGA GAGCTAGAGGGCATCTCAGAGAATGTATCCATGGGAGACACGGTAGTCTCATAATG[A/G]G CTGCTGCACTTGGGTCTGCATGTAACGTACCATGGCAGCAGACTCAATTTGCATGCTACC TGAGCCAACATCATGACTAGCATGCTGTAGTTCTGCTCTTGACTCATCAACAGTATATGCC GGCATAACATAGAAAAGTAATGACACATAAGGTAGTTCGCTGGACAACTAATATGT
>Gm03_40340517	TGAGATTACTGAGGCCAATTCCCATTCCATTAGAGAAAGCAGTATGAGCAGGATATTCTGC ACCAGTTTGAGAGCTAGAGGGCATCTCAGAGAATGTATCCATGGGAGACACGGTAGTCTC ATAATGCGCTGCTGCACTTGGGTCTGCATGTAACGTACCATGGCAGCAGACTCAATT[T/C] GCATGCTACCTGAGCCAACATCATGACTAGCATGCTGTAGTTCTGCTCTTGACTCATCAAC AGTATATGCCGGCATAACATAGAAAAGTAATGACACATAAGGTAGTTCGCTGGACAACTA ATATGTTCTGTAGGATGAATATTTGTGACAAAAACAGCAGGCGGAGGATATTGTAAAT
>Gm03_40377254	AAGTTGCCGCAAGAGAAAAATACATCCTATTTCATATTTTCAGAGTTGCATTTCAAACCAAAT TATAAAATTGAATGATTTTTTAAATATTAATTTTCATGGTGACAATTGCAGGCAGTGCCAAG TGTAGAGCTTTGGGAGCTCGAAGTGGTCTTCCACGTCGGACAGATCCACATGTG[C/T]TATC GTTTTTACACTAGACAGAAACAAGTGGTATTTCCCATCAAAGAAGCTCAAAGCTCTTTTCAT CTCTCTGCTTTTGGAGTTATGTTTACTATGTGGCTCTTTGTACATGTCTTGTGTAAGATGTC CTATAGTTTGTACAGAATCATGAGTAGGGGTGCTCATGATTTGGGTATTTA
>Gm03_40537578	TATTTTTATTATTTTTATTGACGGTGAGTGATGAAAATTGAAACAGTGCAGACGACGATATC ATGGAGCTGCTATGGCACAACGGCCAGGTCGTGGTGACAGTGCAGAACCAACGTTCTCTG AGAAAACCTCCGCCGGTGACCAACTCCCACGACGCCTCTCCCGCCGGTCTTCAAT[A/G]A CGAGAGAGATTGACCGCTGGTGAGAAATTCATCAGCATCTCTTCATGCACGAAGGCG AAATGGCTTCGTGGCTTCACTATCCTATCGACGACGACGAGCCTGCGTTGGATCAAAGCTT CGGCGCCGATTTTCTCTATTCTCCGCCGACGACGGCTAACACAGCAGCTTCATGCAA
>Gm03_40623046	CACTTATATGTTTTTCTCCACCAATTACATCCAATGATCCAAGGAATCTTCTTGCGACTCTT GCTCTATAAGAATACCTATTCTTCTAACACTCGACACACCAACCAACCTAGCTAGTTTT CATCACCTTCCTATGATTGTTTTGTTTTAATTTTCTAACTAACTAACACAGCA[G/T]GAAGCA CTGAGTCTTTGGCCATGACGTTCAACGCGAGCACCATCACAACGGCGTCGGAAGGAGCCT GGCAGGGCGATAATCCCCTGAACCACGCTCTTCTTTGTTGATCGTTCAAACCATCCTCGT AGTCTTCGTGAGCCGCACACTCGCCTTTCTCTCAAACCCTTTCGTCAACCTA
>Gm03_40670148	CCCCTGGAGAAAGTGTGGCGTACAGCGCTAGGGTTTTGAAGCAAAAGATAACAAGGATAT AGACAGCGATAGAGATAGAAAGGAATGAGAAAAAGAAGAGAGAAGCGAACCTTAGAAAGG AAGAGAAGCGTTGGGAATGGCCGCACTTGTTTCGAGTGGCGCTGTGGTTACCCTAGAAGA[ G/C]GATGGGAGCACAAGCGAGACTAGAGTTTGTTCGAGTGCCTATTCTAAAATCTAAATC TGTGGACCTATAATATACTGGGTTGGGCCGATCTCAGTCCACTCTTGGGGTATTGCTATTA GCAATTCAAAGTTTGCTATTGGTAGTGGAATAGTAGTCCCCTTACTCTTATTTTGTGCGG
>Gm03_40685818	AAATCTAGGTCTCGCATCCTTTACATGTAATTTAAAGCTCAAACAAATTCCTCGATGCGGGC TTCTGAGATTTAGACTCTGAATAGCAGGGGAAATCAATCTCTGCTGAACACGGGAAATGAG ATGTTGGCTATAACCTCTGCTCGCAACACTTTGAACTTGAATTTCCGATGAAGAAT[T/C]ATG TCTTATCTAGTACCTGTCAGGCTGTCACAGCAGGGGAAAACATGAGACAAGTCAAACGGGT AATAAAAAATAAAAAATGTGAAAAAGTTTCTCAGTTGCATTGAGAAAACTGTAAAAAAG ATCTCGCAATAAAGAACAGCATAACACTAAGACTAATCACATAACGGAATGTCA
>Gm03_40689982	TCAGGAAAAAACAACAAAAAGTTCTCCAAGCTGGAATGATAAAATTAGCCTATTTTAATTT GAATAAAAAGACATAAAACAAGAAAAATAATTTCAACCTATTTAAAAAAGCAACATAAAATAAT ATAAAAAACCATTAACCTCTTTCTAATAGAAAGGTGAGAATATAAAATTATAC[A/G]AGATT AAAACATTTGATATTGGATTGTGCATGTGCTATAAGAGGCAGGTAAATTTGATATTGCAGGG ATGACATATTTGAAATTAAGCACAGTTTGTAGAATGCATGAGCTAAGAGGCAGCTAAAGT AAAAATGTTTTTTTTTATTCTACATAATATTGATAATGCAGCCTTTTTTCA
>Gm03_40760791	TCTATTGTATGTCAAATAAAGACCTGGCTTCTCTCTACTGAAAGTTACACAGTGTGGAT TTCCCATGCTGACACAGGTCACATGCCAGATAACTCCATCTACAACATAGTTCTGATTTAACA ACAGCATTATCCTTATTTGCATGTAATTTAGTAGGCACATCTGAGGCTTTAAGAA[G/C]TGGT TCCCCATATCAACTCTGACCTGAAACAGATAGCATTATTGTCTATCAAGCTGAAGAAAGAG AACCAATACCAGATAAGAAACAAAAACAAAAATCCCTTCTGACGTTAAACCAATTAAGGCAC CTTAAAGCAACATGGGTGATGTCTTTAAACAATATCACATATTAATAAGT
>Gm03_4078529	GGCCTTACCCATGGATTCTCTTCCCAAATTGCGCCACCCTTCTCCATTTCCCACTCCCTTAT CCTCTTCCCTCTCCTTCCGCTCTTTACCGCCACCCCATTTTCTTCGTGCTGCTTTAGGTTA

1	CCCTCCATCAGAGCCTCCTCCTCACCCCTCCCAAAACGACCCCGCTTTTCGCACCC[G/A]AA GAAACAAGCCCTCCCTCCCCCCTCCTCCAAACCCTAACCTCCTTCTCCTCCCCCTCCT CGAAACCACCTGCATCGTCATCGCCGCCGCCGCTTCTTCTTCATGCGCTTCCACCACAC GCCCGCCGTCATCGCCGCTCCGCTCACTTCCCCCGCCGCGGAAACCGACACCGCCTT
>Gm03_ 4078529 9	CCATGGATTCTCTTCCCAAATTGCGCCACCCCTTCTCCATTTCCCACTCCCTTATCCTCTTCC CTCTCCTTCCGCTCTTTACCGCCACCCCATTTTCTTCTGTCGTCGTTTAGGTTACCCTCCAT CAGAGCCTCCTCCTCACCCCTCCCAAAACGACCCCGCTTTTCGCACCCCAAGAAAC[A/G]AG CCCTCCCTCCCCCCTCCTCCAAACCCTAACCTCCTTCTCCTCCTCCCCCTCCTCGAAACCA CCTGCATCGTCATCGCCGCCGCCGCTTCTTCTTCATGCGCTTCCACCACACGCCCGCCG TCATCGCCGCTCCGCTCACTTCCCCCGCCGCGGAAACCGACACCGCCTTCACCCAGG
>Gm03_ 4087065 3	GAACCCCTTGGTTCTGCGTCCCTGGACTCCCACTACATCCTCCTCCTTTCACCTTCCGCGAC CTCTCCCCCTCCCCCGTTGCCCTCCTTCAACCGCTCCTACCCCGTCACCTCCCCCAACGAAT ACTCCCTCTTCTTCCGCAACTGCGCCCCGAAACCTCCGTCACCATGTCCGTCCACA[A/G] CGAAGCCTACAATCTCGACGCTGACGGTTACGCGACTACCTCTCCGCGGCCAAACCCCT CCTCCCTTCCCTCTACTTTCTTCTTTTAACTACCTCTCCTTCTTCTCCTTCTGGCTCTA CATCTGCTACACCAACAACTCTCCGTTACCGCATTACCTCTTAATGTCTCTCT
>Gm03_ 4088538 2	ACTGGGCAAGCACATTCTACGATGCATGAGGCGTTCCAACAGTTGAAAACCTTTCCCTTTC ACACTTTTCTTTTTCTCTCACTTTCACTTCTCAGTCTCAGGAGAAACCTACGCCCTTCCGCTCATC CTAGAGCTCCACACCCCAACAAACCCCGATTTTTCTCAACACACACTCACACAATC[T/C]GT TTTTCACTCCCTCGATTTTCGCGATCTCGCAGTCGATTCTCCCTTTCTTTTAATTATAAACGC CGTCGTTTGCTTCCCCGCGAAGTAGTTAGGGTTTCTTGTGGCACAATTATTGGATCGCTG CATCGATAATTGACGGGAGAAACAATTTTTTTTTAATCGAGAAATTGTTTTCTC
>Gm03_ 4088633 3	TGTGGAATTAATCAGTTCCACTTTGCGATTGGAGCAGGGTTCTTTGACTGGAGAGAGTGAA GCTGTGAATAAGACTAACAAGCGTGTGGATGAGGATGCTGATATTCAGGGGAAGAGGTGT ATGGTTTTTCGACGGGACCACTGTTGTTAATGGGAATAGCATATGCTTGGTCACTCAGA[A/G] TGGCATGGATACCGAAATCGGGAAGGTGCATATGCAGATACACGTAGCCTCGCAGAGCGA GGAGGATACCCCATGAAGAAGAACTCAATGAGTTTGGAGAGAAGCTGACCTTGATAATT GGACTGATATGTATTTTGGTTTGGCTCATCAATGTTAAGTACTTCTTTCTTGGGAGTA
>Gm03_ 4088902 6	CTCCTAGGGAGGAAGTATACCAAGCAATTGAAGACTGCAGAGATGCTGGAATTCGTGTTAT GGTTATTACTGGAGACAACAAGAACACCGCTGAAGCTATATGCCGTGAAATAGGTGTATTT TCTCCAGATGAAGACATTAGTTCTAAAAGTTTAAACAGGTAGAGATTTTATGGAACGT[G/C]AT GATAAAAAGGCATATCTGAGACAGCATGGCGGGCTTTTGTTTTCAAGGGCTGAACCTAGGC ACAAGCAAGAAATTGTGAGGCTGCTCAAAGAGGAAGGGGAAGTGGTGGCCATGACTGGG GATGGTGTTAATGATGCTCCTGCCTTGAAGCTTGCAGACATTGGTATTGCAATGGGCA
>Gm03_ 4095112 3	ACAAACACGTTTTGAGACGCCGTTTCCACGACCTTTCGATTACGCGCAGCATTGAAGGCG ATTTTTTACTAGCATGGCAGCAATAATGTTGACGACTATGAATATTCCTCTGCACACTTCGG TTTTGGTTAATTCCCGGCGGAAGAGTTGGTGTCTGCTACGCCTCACTTTCGACGCCAG[C/T] TAGGAAGGAGATCCGTCTCGTTTGGCCAGCAAAGGTCGCATGTCCGCTGACACCCTCGA ACTTCTCAAGGTCATTCCAATTTGACTCGATTGCGCACTGTTATCTTCTACATTCTACACTA CTAACTAATCTGCATTCTTTCTTCTGTGTGAAGAATTGTCAATTGTCAAGTGAAGCA
>Gm03_ 4100769 1	CAAAAAAAGAAGCCCACTTAGAAGGACTCTTTAAATATATTTATATTCAACGTATAAAACCTT TGATGCTGTTCTGCCAAACCCCAAACCCTAGTTTTGTCAGTAGCACACTTCACCCATGGCA GTAGCAGCAGCTCGCCCTATCGTCTCCGTCCAAACCCTAGAGGGCGATGCAACCC[G/C]TA CCGTTCTCTTCCCGACGTATGAAAGCCTCCATTCCGGCTGACATCGTCAATTTCTGTTCA CTCCAACATTTCTCGAACAGCCGCCAACCCCTACGCCGTGAGCAAGCGCGCCGGCCACCA AACCTCCGCCGAGTCTGGGGCACCGGCCGCCGCTCTCGGTATCCCGCGAGTCCC
>Gm03_ 4100793 7	CAACATTTCTCGCAACAGCCGCCAACCCCTACGCCGTGAGCAAGCGCGCCGCCACCAAC CTCCGCCGAGTCTGGGGCACCGGCCGCCGCTCTCGGTATCCCGCGAGTCCCCGGCG GTGGTACCCACCGCGCCGCCAGGGCACCTTCGGAACATGTGCCGTGGCGGCCGCGATG T[T/C]TGCTCCACCAAGATCTGGCGCCGCTGGCACCGCAAGATCAACGTTCAACAGAAGC GCCACGCCGTCTCTCGCCATCGCCGCTCCGCAATCCCTCGCTCGTCCAGGCTCGC GGCCACCGCATCGAGTCCGTCCCGGAACCTCCCCCTCGTCTGTCAGCGACACTGTGAAAGC GTCGA
>Gm03_ 4100818 9	CGCCATCGCCGCCTCCGCAATCCCCTCGCTCGTCCAGGCTCGCGGCCACCGCATCGAGT CCGTCCCGGAACTCCCCCTCGTCTGTCAGCGACACTGTGAAAGCGTCGAGAAATCCAAAG AAGCCGTCAAAATCTCCAGAAAATCGGCGCATTACAGACGCGGAGAAAGCTAAACTCA[ A/G]CCACGGAATTCGTCCCGGGAAGGGAAAAATGAGGAACCGCAGGTATATCTCTGGCAA

	GGGTCCTCTTATAGTGACGGAACAGAAGGTGCTAAGGCCATTAAAGCCTTCAGGAACATT CCCGGTGTAGAGGTGGCAAACGTTGATAGGCTCAACCTTCTGAAGCTGGCACCCGGTGGC CA
>Gm03_ 4100825 5	GGAACCCCCCTCGTCGTACGCGACACTGTGCGAAAGCGTCGAGAAATCCAAAGAAGCCGT CAAAATTCTCCAGAAAATCGGCGCATTACAGACGCGGAGAAAGCTAAACTCAGCCACGG AATTCGTCCCGGAAGGGAAAAATGAGGAACCGCAGGTATATCTCTGGCAAGGGTCCTC[T /C]TATAGTGACGGAACAGAAGGTGCTAAGGCCATTAAAGCCTTCAGGAACATTCCCGGTG TAGAGGTGGCAAACGTTGATAGGCTCAACCTTCTGAAGCTGGCACCCGGTGGCCACCTTG GAAGGTTCAATTATTTGGACCAAGTCCGCATTTGAGAAATTGGACTCTATCTATGGATCCTT
>Gm03_ 4117022 9	TCAGAGATTCTGTGTTACATAGCCAATCCTCTCACCAAGTGCAAGTAGTTCCTGGTCAA ACCAGTCATCAAAATAATTAATAAATCTGAGTGCATAATTAATTGCAACGCTAGAAACAG TAATAAATTAATAATCACCTCATAACTCATGTTATCTACGTCCATCCTCATGTCTC[T/C]GTGC TGATCTAGCATGTTCTTGGAAACCATACTGATGCACGCTCAACAACCATGAAACCCTGAA AGCCAACAAAAAATACAAAGAAAAAAGTAAATGAGTCTCAGCCAGTGGTAGTAATGCCAG AAAGAACAATTTTGATGTTCTTAATGAACAGCGCAACTAGTTCAAAAGAATATA
>Gm03_ 4118456 4	TTTAACAAATGGATCAAGCTGTCCACTGATCTGTAACCGTGGTGGACTCGACAAGACAACA ACAATATTCAGAACATTAACAAGACCACACCACACCATGAGCGTCATCGACATTCTACCA GAGTTGATTCCATTTGCAAAAAGTACGACAAATACGACGTCCAAAGCCAAAGGGACT[T/C]T AATCTCTCCTCCGACGATGCATTGCGCAAACCTCTACGCCTCCGTGACGCCGACATTGAGG CCTTACTTCAGGTCACACTTCCCTTCACTTTTTTTTCAAATTCATTACTAAATGAGGAAATT TGTGCCCTAATTCGTGAGAATTATCGATTTTTTTTATTCATCTCCCAAACATGTG
>Gm03_ 4122368 1	TATCAGCTATTTGTGAGGCTTTGAGGAGTTTGATGAAGGTTGTGAATTGTCCTGCTTTCTCA AGCACTTGGGTGATGTTGGTGGGTCCCGCTGGTGCCGGTGCTACTTGGGCTGAAATTGTT TGGATGAAGATGTTGAGGAAGAGGAAAGGGAGGAAATTGAATAGCTTTGTCAATTGGT[C/T]T TCCTTGTGAGTGAGGAAATTAAGAAGAGAAGACTAAGATGTTTGTACAAGTGGAAAGGAAG AGATGTGGTGTGTAATATATATTAAGAAGAGGGTGCATGGAAATTTGATTGTTTAAAGGT TAGGTGAAAGGTTGCAATTAGTATAAGGTAGCAAGAGGCCCTCATGCAATCAATTT
>Gm03_ 4122828 4	TAGGCCTTGGTTTGAGTTGTTGAGCTGTGAATTGATTCTATCAGCTATTTGTGAAGCTTTGA GGAGTTTAATGAAGGTTGTGAATTGTCCTGCTTTCTCTAGAACTTGGGTGATGTTGGTGGG GCCCGCTGGTGCTGCTGGTGTGTTGGGCTGAAATTGTTGGATGAAAAGGAAAAGGA[T/C]A AAATGGATTAGCTTTGCCATTTGTGTGAGTGAGGAAGCTAAGAAGACTAAGATTTTTGTAC AAGTGAAGAAGGGGATGTGGTGTGTTGTTATATTATATGGTGTAAAGGTTTGAGGGT GAGTGGAATTAGATTGGTATAAAAAAGTTAGGTGAAAGGTTGCAATTTGTATAAGG
>Gm03_ 4122832 1	CTATCAGCTATTTGTGAAGCTTTGAGGAGTTAATGAAGGTTGTGAATTGTCCTGCTTTCTC TAGAACTTGGGTGATGTTGGTGGGGCCCGCTGGTGCTGCTGGTGTGTTGGGCTGAAATTGT TTGGATGAAAAGGAAAAGGAGAAAATGATTAGCTTTGCCATTTGTGTGAGTGAGGA[G/A] GCTAAGAAGACTAAGATTTTTGTCACAAGTGGAAGAAGGGGATGTGGTGTGTGATTTAT ATTATATGGTGTAAAGGTTTGAGGGTGAAGTGAATAGATTGGTATAAAAAAGTTAGGTGAA AGGTTGCAATTTGTATAAGGTGGCAAGAGGCCCTCATGCAATCAATTTTTCTTTCTA
>Gm03_ 4123079 7	ATTGGCAGTTCCTTCCATTCCCATGCCATCGCCAGCGCATATCAAATCGAACAGCCGCAAA CCCTTCTCTGAACCTCTGATCCGCACTGGCACTGAAGCCATTGCTGTACCTTCCAAACCCT AGCGTAGCCACGCGCTGAATTGCGATACGATGATCGTGCGCACATGAGGCGGCACGG[T/C] JCCGGCACGATGATGAACACAGCCAACGGCACGATGATCTCACCCTCGTCTTCGTGCGGAA CGGCGGCGCAGTCGCCGGGACTGAAGACCTATTTCAAACCCCCGAGGGAAGATATAAGC TTCAATACGAGAAGACTCACCTTCCGGTCTTCTTCATTACGCCCATGGCAAAACCGTTA
>Gm03_ 4123154 7	TGACAAAATGGAATTGTCTTAGCATATGAGTGAATTGTTGGTTTGATTGAATGAGCAGGTAA CCCTGGCACATCTCAAGGACAAGCCTGCGCCTTACCCCCAACCGCTTCGTATCGTCAA GTTTCAGTGCCAGCAGTGGGGTACGGTCCGGCGCAGCTCGATTGCTGGGAGGAAGCA[T/A] JTGGTAGCCGGGCACTTAGTTTTGTTGGAGGCAATGGTGGCAGCAAGAGTAATGGAGGAAA TACTAGGATTAGTTCCATTGGTGCTTCGAGTACAAGTAGTGCAATGGCGAATCCGAATTTT GACGGGAAAGGAACATACTTGATATTCAATGTCGGGGATGCAATTTTCATTAGTGATT
>Gm03_ 4123325 2	GTATACGAAAAGGCAAGTCTCAATCATCAAAAGAGAACTTACTTTATGGTTCATTTATGAA TATCTTTCTTATGATGAGTTAAGTAGATATTTACCTCTTCTTTCCTATTTCAGAACAAAGATG GTGCTGGTGATTCTTCTTTCCAGTTGTAAAGGATCAAACCTCAATTTTCGGTTG[A/G]CCATG CACGTTACAGTAAGGTATTGAATTTGATGGCCTTTTGATTTCTTGACAACCTTATTGATATGCA ACTTCTATTTTACCTTTTTCTCTCTTTTGGCCTATCTATTTGATTTTACTGTAGTTAAATTT AGTTTTTATCACTTATTAGTTATGGTTGCTAAGTTTTTGTAGAATTTT

>Gm03_4123433 8	GATGATTTAGTTCAAGTTTGGAGCATGGAAGATCGCAAGGTTGTGGCATGGGGCGAGGGA CATAACTCATGGGTAAGTTGCCTGACAATTTTATGTTGTGTGCTTGTAGCAAACCTTTGCTAC CGTAGGGTTAATGTAGTGTTTGTATCAAATGCCATGATCAGGTCAAGTGGAGTTGCTT[G/A]T GATTCATATTGGACATCACCAAATCAAATGACAATGGAGAGACTGTTACATATCGTTTTGG TTCTGTTGGTCAGGTGCTCAGAAAAATGTTTCATTATAATTTTTACTTTTTCTTTGTGTAGCT TTTCATTCTAAGTAAACTCATTGGTATACCTAACATGGATCTGTGGTAGCTTT
>Gm03_4125534 6	GCCGGCAGGGAAGGACAATTGTATCTCGGGGCTCTGACAAGCAAATAACACATTTTCCTTG GTCATTCTCCGTTGAATTTCTAATACCATATATCTCCTGCAGCTCATACCTCATTCCATTAC CCACAAGATCTGCTTGACAACTTTCACCCGGAAGTCTCCTTTCTCCTTCTCAAAC[C/A]TCGT TTGCGTAATCTGAGAGTTTGTGTTACTAGATGTTGGAGATCTATCTGACTCATCATGATTAC CAGGGGATGCGTCTGCCTTTACTGCTAGAGGGTAGACATCCTTTTCCACCCACCTTTCCAA TAACTCAGACTCTTCAAACACTGAAAAATCAATACCAGTTCAGCTGCCTGTC
>Gm03_4127578 8	GAAAAGGAAATCGTGAGAAACGGGAGCGAAAAACGAGTTGATTATAGTAAGAAAGCGCGTG AAATCCGAAAAGGGCTGCATTGGCGTGAAAAATGGTAGATCTAAAAGTAAAGGAGAAAAACCG AAATAAAGAGATTGGAAGGTACCGTAGCAATCGGGTCTGTGTTAGATTCAGAGAAAAC[T/C] AGACACAGTGTCTTAACGAAAACTGGAGGACTGCTTCGCTGATTTGCTTTTCGCTCTTCGC CCAATCAAATCAACACAACCCACCTATTTCTTTAATAATAATAAATCACACATATATAATTT AATTTTACTCACAACCATATTCTTTTCGCAAAGTGTGGACATAAATAATAATGACA
>Gm03_4197769 7	TAACACAGTTCTCCTTTCTCTGAATTTTTCGTTAGAGAATGGGGAAGAGTAGCGGAAGAGA CCGAAGACGAAGAAGATCACATGACTCTGAATCCGACTCACCTTCCGATTCCGACTCCGAC AGACGCCACCGTAGCAATAGTCGCCGGTCGCGGCGGGACTCCGATGGCGACAGGAAG[C/ G]AGAAGTCCTCGAAGTCTGAATCCGAGTCGCCCTTCCGATTCCGACTCCGACATACGCCAC CGTAGTAGCAGCCGCCGGTCTGTGGCGGGACTCCGACGGCGATAGGAAGAAGTCCTCGAG GAGCATCACAGAAGAGGAAATTACAAAGTACATGGCCAGAAAGGCCAATCGAAGGTCTC AC
>Gm03_4217910 9	GGCAGAGCCACTGCAACAGGAAGCAGAGTTGCGCGCTACTCCGCGACCCTACCCTATCAA TTTTCGTACTTCTTTTCGTAATTGCAGCGAAGAATGGCGAAGAGGAAGGAGCGGATTGAGA ACCCGGAGCCATTTCGATCCGTACGGCGCCGACCCCGCGAAGACGAAGAAGCGTTTCGAA[T/ C]CCGCCGAAGCGGCACCAGCAGGAAGAGCAGTTTCATTGCTCCCAAACCTCAGCTCCAAGAT CATGAAGCAGGCGCTGATCCAGCAGAAGGAGGAAGAGAAGGAGGAAACGCACGAGAACA ACGCCGCCGCGAATTTATTCGAAGAGTTCCGAACCTTCGAGGAAGACGGCGGTGACGACA TC
>Gm03_4222412 5	GTTGGAACCGGACGGGAATCTGAAGGGTTATTACTGGGACGGTTCAGGTGGATGCTTAA CTACCAGGCGATTACTGAAGCCTGCGAGCTCCCCGCTCCTGCGGTTTCGTACGGTTTGTG CACGCCGGGTGGGTGAGGATGCTCCTGTTTAGATAACCGAACCCGGTTCGAACCCGGTG[ T/C]GTGTTTTAATGATGCTTCTTCTGGAGACGCAGACTTGTGTAGTAGGAAGGAATTGGTG GAAAAAGTAGTTACTGGGTGCTAAGAAGAACCAGGAGTGGAGGCAGCGCATAAGGAGCTAC TAAGGCACGTAACGACGTCGTCCTGCGGAGTGTGAGGGATTGTGCCAGAACAACCTGTA G
>Gm03_4233476 9	TAAAGAACACAAATAGGAAAAATACAAACAAGACAAACACAACACACAGGTCCATGCCAGA GCATGGTGCACACTCAGACACAAGTTTAGAGAAGTTCCATACTGAGGTGAAGTACTCAGCA GTGTGATTGAGCCGACTGAAGCCACCGAACAAGGCATCATCTGAATCCAAGACAATC[A/G] GGAAAAAGGAAATGCATAAACCCATTATTAACATTCTCAAACCTTAGGAAAAGAAAAA GGAAATAATATTGGTATTGGTCATAATGTTGGAAGCATTAGCTTCCAGTGATGGTATCGGTA GTACTATCGTATCAAAAGTACACAGTAGGACCAGACCATCCATAGATGGAATTGGA
>Gm03_4458162 0	CGTAAACAAGAAAGGTAAAGGTAAGTCTGTAGATAGTAGGGGGAGTAACTCCAGAGAGAA ATTAGCAAATAACTTTTTAAGCACTGTACATCAGCTACAGGCAAATTATAAATCTTCAGAAG AGAAAGCGGAGGTGCTATCAAGGGGTGATTACCGCAGTGAAGGAGGCAAATTAATAAA[A/T] TGAGCAGCGGATTGATATGAATTCTGGTAGTCAACCTACGATGAAGCTTGGTGGGAAGACT GAAACATCAAACGACAGTTTATCTAATCAAAGTAAAGACGATGGGGGTGGTGGGAACAAGC AACGCAAGAAAACTTCTAAGTTCTGAGAGTACGCCTTGGTGATGGTTCTGTTTCAGC
>Gm03_4477915 9	CTTGCTTTTTGTTTCGATTGAGACTATATGTTATAGCGTATTAGTATAATATAAGCCTTTTTGT CTTCCCACTATGATTGGTGGTTGTATCATTTTTGTTGTAGATTTTAGGTTGTTTTTCCCACT AGGAGTGTTGAGGTTTCAGGTGGAAGTTCGTTTTTGTAGTGATGGTCACTTTT[C/G]TTCTG TACAGTAGTAGTGTGTTGAATAAAATGAAGAGGCCGTGATTGGAGAAAGTAAAGTAGTGTGT GCATAGCTAGTGAAAAATATGTAGTACCAATGTGAACGCCCAAAGAAAAAACACTGGTGAAA TTTTGGATTCATTTTAGCATAAGTTTGGTTATGATTTTATTCATCGCAAATT

>Gm03_45034077	AAACATTTTTATATGTTAACACCATTTTTCTTTCCCGGATGTCACACAGATTACATTTGGTGC CAAGAAGCAAAATGTTGCAAAGAAGCAAAATGTTCCGATCTCCTCTATATTTAGTAACGATA GCGATGAAGACTAATGATGGAACATTACTCGTGAAGTGACATATCAGCGTGATG[G/A]TGAA TCTCCTGTGTGGTTTGGTGTGGTTTGGTTTGTATTAGCTGTTTTATTTGATAATTGTTGCCAT CCTCGGTTCAAGAGGAAGAAAACATTTGGGATTATATGATAGTTTGGAGTTTGGTAGACCA AAAATGTTACCTGCTTATCTGTGAGGTAATGTTATGTTTGAAGGATTTTAT
>Gm03_45105650	GAAGCAAACATGGGGCGACTTTGAAGATAGAAGGATGCGACCATGGACTCCATTATCGGT GGTGAAGAGCGGGAGCAAGCGGATAAGAAAGAGAGCATCGTGAACGTGGTTCAGTCTTGG CTCGTGTGATCTATAATGGCTTAGCCTTGACGGCGGGCGGTGTGTTTTGCCGTCTCCTC[G/ T]ACTTTTCTATCTTCGTGACGGTGGCGACGATTGGTTTAATTAGTTTTTTTTTAAAGTAG GGTTAAAATGGCCTTCTCATATATAATTTATTTTGATATATTTTTTAAACCAACTACTTCAT TTTAATATATTTTTTAAACCAATTACTATCTTATAATATGAACCGTCCAATTATT
>Gm03_45392239	CACATTAATTGATTTTTCAAGGGTAACACTAGTATTTTTGTGCTCTTGTAATTTTCTGT AGGATGGAGGACCTAATCCCATTTTTATTCCAATACGTTTCAAGTCGTGAGAAGCCTGGGG GAGATGTGTTATTTGTAGCTCACAATGCTCGCTGTTTGATGTGCCCTTCATCAT[T/C]AGTG AGTTACGGCGTTATTCCATAAAGATTCTCCTAATTGGTTGTTTTGGACACCCTTCCTTTA GCGCGGGAATAATGAAGTCCGGAGGTCTCTTCTCTCCTCTCTTAAACACATATGC ATGCACGCACACGCACACACACACAAATACTTATTGTTTCCATGTGAAAACCT
>Gm03_45503654	CTAAATACACAAATAGAAAATGTTTTGAAAAGGAGCAGGGAGTGCAACATTATAAAGAGTG GCCACTCAATAGTCAATAAGGGGAATCAAAAAATCAAATGCTAGTTTTTTGTAATACACGTC AATGTAAAACATGGGAGCCACGATATGTTATCAGAACAGAAATATCATAAAAACTG[T/C]CA CCGACGCCTAGAGCAAACAGTTTAATTTGAGAAGTGAACAATTGAAAAAAAAAATAGAACA AAGTTATATTATTTCTTAAACTCTAAATCCTCTGCTGTTTCTCCTCCTCTAGCCCTCTCG AACTTTCTCCCTTTGCTCGTACATAGGGCTTGGTATGGCTGTGTGGCACACC
>Gm04_44751336	CCGTTAAAACCTCAACTGCAACTGCAGCAGCAAACACGTAATTTCTTCCATGCTCA ACGTAGCAGAACGCGAAGAAGCACTACCAGATTCCACTCATCCAACGCTGCTTTAGTTCT TCCGCCTTTCAAATGCGTCCGCGTTTCGCCGCGCGCGCGTGTGCCAGGTCGCCAGC[A/C ]TCGACTTCGAATCCGCCGCCAAAGCCGGAGAGGATCGTCTCTAAGGTATCATCGCTAC TACTCACTATTTCTCGTGTTCCTCCAAATTAGCCCTCGTTCGTAGCTACTAGTACAAGCTC GTTTGGTTCCGGTTTTGGTTTTAGGTGCCGGTGCCTAATATAAGGAACCTCTGCATCA
>Gm04_44761104	CTCTTTTAAATTCGCTTATTTTACTTTTCTTCACTACATTTAACTGGTTTAAATTTGTCCGGC GATTGAGCTCATCCAGCATCAACAACCTCAACTTTACCCACACTTATCTACCATTCTTCAAA TTCGAAGCCTCCTAGCTCGGAACCTGGAGTGTTCCGTTACAACACTGCTATCGT[T/A]AAGGC AATGCCCCAGCGGACTGGTTAGCTAAGCATGGTGCTTCTAACGCACTTTTCTCTCCGTCC AGCCTCACCTTGCTGCTTTTGGCTGATTCCCTCGGAGCTTCCATTGCTAGAACCTAGTTCTG TTTTCTGCTTGCTGCTTGTATTCTTACATGATAAAAAAAAAAACTCGTTT
>Gm04_44772822	TTGACCTAGGCTTGTTCCGCAAGGAGTCTCGGGTAGCGGGATTCCGTCGTATAACT CATCGTCCGCAGTGGTAGAAGCCTTCCCGTTGGTTCTAGAAGGTTCTCGAAGATTCCTCAC CATCCTTTGGGGCGCGCGGTGAAAACAGCGGAGTCGTGTCGTAGCTGCCAGGCTTA[C/ G]TCTGATGAGCAGAATCGGCGCGCTTGACAAACACTCGGTATCCTCGCGGAGCGCGA GCAGAGACCTCCCATCACTTGCGACAACCAACCAAGAATTCAATCTTGAATACGAAGCC CCTGCCATCTAGTAGTAGTAGAGCAGTAAGAGTTACCGCTCTAAGATCGCAAGGCTCGTT
>Gm04_44787477	AAAAGATAGAACCCACAAAAAATATGTATAGTTATATATAGAGAGAGAGAGGGCCTTG GTTGGTGGTGCAAGGCATAAAGAGGAGAATTGGTGGGGATAGGCGAAGGAGGTGGCGAC TGAGTAGTGAATAGGGTTTTTATTGATTGATTGATAGCAATAATTGGTGTTAGCGAAGCTTG[G/A ]GAATTGAGATTTATGGGTCGCAATCGCAGTCGCAGTCGAGTGTGTACGCTCTCCTTCC AATTCCCGCAGAAGAAGGTACTCTCCATCCCCTGTTTCTCATCGCCACTCGAGAACCTCCA GGCGCCGCTCTCCTTCCACAAAGCGCCGTGCAAGGCACAGAACCCTTCTCCTCCCCCTCT
>Gm04_44787569	GGTGGGGATAGGCGAAGGAGGTGGCGACTGAGTAGTGAATAGGGTTTTTATTGATTGAT AGCAATAATTGGTGTTAGCGAAGCTTGAGAATTGAGATTTATGGGTCGCAATCGCAGTCGC AGTCGCAGTGTGTACGCTCTCCTTCCAATTCCCGCAGAAGAAGGTACTCTCCATCCC[A/C] TGTTTTCTCATCGCCACTCGAGAACTTCCAGGCGCCGCTCTCCTTCCACAAAGCGCCGTG AAGGCACAGAACCTCTTCTCCCCCTCTCTTTCACGCAGCCCCACTCCCAAACCCAAGAAA GATCAAAACAAAAGGTAACCAATCATAACCATTCTTTATTTGCCTGGATTTTATCTTA
>Gm04_45033967	ATGTCATACAAAAAGTGTGCAGAAATAACAATAGAATAGTTAGTGGATTGTTCAAAGATAGA CATTGACTGTAAGATGAAAAGTTGAAACTTTCCAAATGAAAGAAAGAAAGGGAAGAGAGT ATTATTCAAATACCTGATCTTGGGGGAACCTCACTTGCACAGGAGAGCAGTGATCA[A/C]GA

	TTGTCACCGACTCCAACCTGTCCAACTACAATTGAAGGCAAAATAGGTAAGAATTTTCCAA ACACATGTCAATAGTAAAAGACTTGTGATTGTACAGAAAGAGAATAAAGTCTTAATACCAAT TTTTTTTTTATATGAAGTAGAATATATTTAGTTCTCTGCACACATGAATGTGCA
>Gm04_ 4503478 7	AAATGCCAATGCATCCAAATAATACAAGAAATTGCTAGCTAGACACCAACTCAGCATAAAAT TAAAGGACTACCTGAGAGATTAACCTATCACTCAAGGCTTGAAGCTTCTGAGGCACAAGAG AATCCTCAAAATTTCCATGTCCTAGCTGGCCATATTTGCTCCACCCGATGTGTAT[A/C]TGC CACCAGAAGATGAAACAGAAATTGTATGCCGCCAACCACAAGCAACCATGACCATCTTGTC ACACTGCAACATTAAACAAGCAAGAATTAGGCAAGGGGAGGAGGGAGAGCAAAAATCAAA AGTTCTCCATCCATTACTATGCATGTACACCAATCACACATGAATTTTAAACAACAT
>Gm04_ 4506150 9	AGAAAGAGCTGTTCAACTTTGTGATGAGTCACTTGGATGCAGCCACACAAGATGAAGAAGG GTCTGCACCTGTTTTGTTTCACCCATTTGGGAGCTACCTTTTTGTTTCATCAGATGGAGCAG TGTGTGTGGATTGTGCAACTTCTGTTATAAAGATGTTGCATTTGATGTTTGCATCA[A/T]TTT CCCCAGCTTGTAGTATTGAGGACCATCTAAAGTATGGAGACTATGTTAAGAATTTGTCATTA CAATTCCTGAACCAGAACAATTCGTCAGGGGAATATTCCTGATTCAAGCTATGAAGCAG GACTTGAATTGTCGGTTCAATCTTCAGGTTTGGGTAACCAGGTAAATGCACTAT
>Gm04_ 4509041 9	CCAACCCCCCTCCCCAATAAAAAATAGGAAAAGAAATTAGCTAACCCTAATGACTGATGA TGTAACAGTTATGGAGGAACCGTTGGCAGCTTGAAGAACATTGCGACTAGAAACATGTCA ATTGCTTTGCAAACCAAGGCATACAATCTGACTGTTTTTGCCTGTATTCTTACAAAT[G/A]CT GTTGCAAAATTGGTGCAGTTACTGGAGTTTGATTGGTTTTACTATTTGCAGAGACATTGAAT TTGTACAGATTTGCATGTATAATAATACATCAGTTTTTAAACAAGAATTTGACTGAGAAAATCA CACGCCCCGTCCCTCTTGCAACTACTTTCACATTATTTACGTTAGATACTTCC
>Gm04_ 4538076 4	AAAAATAGATTGCACACCTCAATTCTGTTGTTTTCTTTCACAGCCAATGTGTCATCATCTTCA AAGAAACAATCAAGCAAAGATAGAATGTCATTGTCAACTTGATTATTTGGAGCTGAAGGATG TGCTGAGCAAGAGGTCATCAATTCTGGTACACAAGATACAGAGCCAACCTTGATT[G/C]CTTG GTGCTGAAGGATGGTCTGAGAATTGAGGGCATCAATTCTGATGCAGATGACACAGAGCCAA CTTGACAGTTTGGTCTGAAGGATGTGCCGAGCAAGATGGCATGAATTCTGAAACACGTGA AACAGAGCCAACCTTGACTGTTTGGTGTGCTGAAGGATGTGCCGAGCAAGAGGGCATG
>Gm04_ 4538079 0	GTTGTTTTCTTTCACAGCCAATGTGTCATCATCTTCAAAGAAACAATCAAGCAAAGATAGAA TGTCATTGTCAACTTGATTATTTGGAGCTGAAGGATGTGCTGAGCAAGAGGTCATCAATTCT GGTACACAAGATACAGAGCCAACCTTGATTGCTTGGTGTGCTGAAGGATGGTCTGAGA[G/A]TG AGGGCATCAATTCTGATGCAGATGACACAGAGCCAACCTTGACAGTTTGGTGTGCTGAAGGATG TGCCGAGCAAGATGGCATGAATTCTGAAACACGTGAAACAGAGCCAACCTTGACTGTTTGGT GCTGAAGGATGTGCCGAGCAAGAGGGCATGAATTCTGAAACACATGAAACAGAGCC
>Gm04_ 4538083 4	AATCAAGCAAAGATAGAATGTCATTGTCAACTTGATTATTTGGAGCTGAAGGATGTGCTGA GCAAGAGGTCATCAATTCTGGTACACAAGATACAGAGCCAACCTTGATTGCTTGGTGTGAA GGATGGTCTGAGAATGAGGGCATCAATTCTGATGCAGATGACACAGAGCCAACCTTGA[T/C] AGTTTGGTGTGCTGAAGGATGTGCCGAGCAAGATGGCATGAATTCTGAAACACGTGAAACAG AGCCAACCTTGACTGTTTGGTGTGCTGAAGGATGTGCCGAGCAAGAGGGCATGAATTCTGAA ACATGAAACAGAGCCAACCTTGATTGTTTGATGGCAAATGATGTGGTTCCGAGCAAGAGG
>Gm04_ 4538088 5	GATGTGCTGAGCAAGAGGTCATCAATTCTGGTACACAAGATACAGAGCCAACCTTGATTGCT TGGTGTGCTGAAGGATGGTCTGAGAATGAGGGCATCAATTCTGATGCAGATGACACAGAGCC AACTTGACAGTTTGGTGTGCTGAAGGATGTGCCGAGCAAGATGGCATGAATTCTGAAACA[C/G] JGTGAAACAGAGCCAACCTTGACTGTTTGGTGTGCTGAAGGATGTGCCGAGCAAGAGGGCATGA ATTCTGAAACACATGAAACAGAGCCAACCTTGATTGTTTGATGGCAAATGATGTGGTTCCGA GCAAGAGGGAATCAATTCTGATACACATGATACAGTGCCAACCTGGATTGCTAGGTGCTG
>Gm04_ 4538116 2	TGATGGCAAATGATGTGGTTCGAGCAAGAGGGAATCAATTCTGATACACATGATACAGTG CCAATCGATTGCTAGGTGCTGAAGGATGTGTCGAGCAAGATGGCATCAATTCTGATACAC ATGATACAGAGGTCAATCCAGCACATCCACTTGTAGAGGGATGGATGTAATTAGCAA[G/C]T GAAATATGGCTTTCACTAGGTTGTATAGGAACAGGTGCACAAACAGCAGCAACAGGGGCA GATTGTACAGAATCAATTTCTCATCTGTTTCCCACTCTTCTCTCATATGGTTTACCATAT TGTGCACCATTCCTAGGACCAGGACCCTCTTTTGGTACACCCTACAAATCACATA
>Gm04_ 4538512 8	CTAACTCAGCATTTGGGAGGCTAATGCAATTTCTCCAGGTAGCAGAGCTCCATTCACAAGA AACAGGGATGGAAAAATCAAAAAGACTCGGCTCAAGTAAATTTAGCCTCAGACTGTTGAAG TACTTTTGCAGATCAGAATCTTCTCACACTACTCCAGGAGAATAGACAATGCACAC[G/C]G CCACCTTTTAAATCTTACTTCTCACAAAACTCCAAAGCCATAACACACAGTGAACCAACCA TCTTCAAATGACTTGTAAAGGCAAAATACTAATTGATGAACAATCACCGTGCAAGCTTAGTC AAAAGGGAACAAGCAAACGATCACCATTGACAAACTATTTCACAATATCCCATT



>Gm04_45580521	CCAGCTAGCATGAAGAGCAAGATTTGCCAATAGGAGCGAGGCAATCTCACCTTATCCACACGCCACCACAACCTTAAATTACATCCATCATCCATAGCAACAAGCCAAAGAAGAGAAAAATAAATATGAATAACAACCCACAAAACCTTGGGTCTTGCAAATTAAGTTCATTTTCCTC[G/A]CTTCACTCTCGTAGACATCACAAACCTTCCCTCCATTTCATCCACGCATGGCTGCCATGAACCTAGCGTATTGGCCTGTAGCTATGCCATATCAGGTGCAGCGTGTTCGGAGCTCAATGGGAAGGTCACTTCCGTGGCCTATGTTGCATGCTCTGGCTACAAGTTGCCATTGATCAAA
>Gm04_45580672	GGTCTTGCAAATTAAGTTCATTTTCCTCCCTTTCACTCTCGTAGACATCACAAACCTTCCCTCCATTCATCCCACGCATGGCTGCCATGAACCTTAGCGTATTGGCCTGTAGCTATGCCATATCAGGTGCAGCGTGTTCGGAGCTCAATGGGAAGGTCACTTCCGTGGCCTATGTTGCA[T/T]GCTCTGGCTACAAGTTGCCATTGATCAAAGCTGAGGCCAGAGTTCCCTAAAGCCAAAGAATCTGAAGGAGGAAGGAGAGGTGCCCTTGTCTCTTGGCAGCTACCCTTTTACCTCTGCAGCTTCCAATTCCTTCTGCCAATGCTGGGATCATTGAAGATTACCTTGAGAGGAGCAAGGCCA
>Gm04_45581542	TGTACTTAATTTTGCCTTCATAATGTAGATAAGTTTATGTGCTATAGGAATTGAATGACAAGAAGAGGTTAGCTACTACTGGAGCAAACCTTGGCAAGAGCATACACCGTGAATTCGGTAGTTGCAAGTTCCCTGAGAACTTCACTGGTTGCCAAGATCTTGCCAAACAAAAGGTCTAG[G/T]TCCACCTCTGATATTTCACTTGCTTTCCCAAATCCTAACTCTTGCACTTGCTTTTCTAATTCAGTAATTCCTCATGCTGTTGTTCTTTTTCTCTTTTTTTTTCTCCTTTTCGTTTCTGGGTTTTATCTACGCAGAAAAGTGCCATTCTTAGTGACGACTTGGAGCTTGAATGTGA
>Gm04_45581810	TTTTCTTCTTTTTTTTTCTCCTTTTCGTTTCTGGGTTTTATCTACGCAGAAAGTGCCATTCCCTTAGTGACGACTTGGAGCTTGAATGTGAAGGAAAGGACAAATACAAGTGCGGTTCTAATGTTTTCTGGAATGGTGAAAGGAGGCAACATTTTATGCCTAACTGCGAACTTTTCG[C/A]CTTGCAATGTACCTAACTGCTTGAATATATATCATATTTTCTGCATTGATTTATTATTATATTGTTGCTGTTGGATTTTATTTGTTGAAAAAATGTGTGCATTAGGCGGCTATTACTCTAGATTCGTTGAAATTACATTTTGCGATATGTTTGAATTTCCCAAATTCATGTTAGCTT
>Gm04_45594453	AGGTAATAAAAAAAGTTAAACAGGAAACTTTTTTCGCTTTTTCAATAATTGGTTGGAGACAAGAACTATTTTTTGTGGAAGACATCTTAGACATGTATAGTTTAGTTTATTTTCAATTTTTCTTAACAGGAAACAATCATTAGTAATTACTCTCATAAAATTGGTGATGCCAGCAACCA[T/C]ATTTTGTGTTGACTGCGATGAGCAGTAGTACATTAGGCCTCATTGAGGTTTATGACAATCATGGCCGTGCTTACCCTTGCAAAGGAAATGAACATGAGAAAATCTTAATCTTATAGATCCCTAGAAAACTTAATAAACTCATCTACAAAATCATAGAAATGGACTCATAACTTCGTAA
>Gm04_45597626	CACACCAAATGGGCGGAAAAAAAACAAATTTTCATGCAGTGAACAAATGGGTACATGAAAATAAAACTTTTTTAAAGGTATTGCGAAAAAAAACCGAGAGAGAGAGTACCCGAAAGAGAGCATCTGACGGTAAAGACGGAAACTTTGGCGGCAGCATGTGGCGTATTGGAA[C/T]TGAAGCGGTTGTTGGTGAGGTCAATTGAGAGACAGAGCGAAGGACAAAGCTTTTGTCTCCATTGATGCCTCACAGAAGAAGCAGTTACAGATATGTCTGAGGGAACCACTCCAGTCGCAGTACCTAGGTTCCGCCACCTAGGTTTTTATTTTTGCAATTTTCGTTTCACTGGGCTCTTT
>Gm05_38251772	TGCTAATATTTCTTTGATACCCTCTGTAAACCCAGATCTGTCTGGTAAGGTTGATTTGTGGCCAAATGAAGATGGTAAACTTGAGCAGCTTCACTCAGATGAAGCATATGAGATGATTCAAAAACACTTGAGTCTCATTCTCGGTAAACAAAGCTGGAGATTTAACATCAGTAGCTGAAA[A/C]CAGCAAATTCAGGGTAGGGCAGGTCTATGCAGCATCAGTGATGTATGGTTATTTTCTTAAGCGAGTTGACCAAAGGTTCCAGTTGGAGAAGACAATGAAAGTTCTTCCAAATGCAACAGAAGAAAGAAATGGTGTTTCATCGAAATACGATGGACAATGCAAGACCCAGTATTGAACAGGA
>Gm05_38252318	CACAGAGGCATTGTTTGGAAGACCTGAAATTGTTGTAACACCTGAGGGGGCAGTCTCTAAGGATGAAAACATCAAAATTAGCTTTGGTGGTTTAAAGAAGCTTGTTTTAGAGGCTGTGACTTTTGGTTCTTTTCTCTGGGATGTTGAGAGCTACGTGGACTCAAGGTACCATTTTGTGT[T/A]TAACTGAGTAAAGCTTTGTATGTCTATCTTCTTTAGAGACCACAGGTGGGAGCCTTGTGCATTCA GTTGCTCTTTTGAATTTCTGTTTGTCTCCCATACATCACAAAGATACAGTTTCATAGTGTGTTGGTACTGACTGATGAGTTTAAAGTCCAAATATTGTTTTCACTAATTCCTA
>Gm05_38278658	AGGTTCCCTCCCTTCAGATCTGGCCTCCTGCGTCAACCTCCGCAACCTCTACATTCAGCGCAACTTGCTCACCGGCCAAATTCACCCTTTTTGTTCATTTGCCGGACCTCGTTCGCTTGAA CATGGGCTTCAACAACCTTCTCCGGCCCCCTTTCCGTCCGCCTTCAACAACCTTACCC[C/G]TTGAAAACCTTTGTTTCTCGAAAACAACCAGCTCTCCGGCCCAATCCCCGACTTGAACAAACTCACCTCGACCAGTTCAACGTCTCCGACAACCTCCTCAACGCTCTGTCCCTCTTAAGCTTCAGACATTCCCTCAGGACTCTTTTCTAGGTAACCTCCTCTGTGGCCGACCACTCTC
>Gm05_38279366	GGAAGGGAACGCGAAGAAGTTGGTGTTTTTGGGAATGCGGCGAGGGCGTTTGATTTGGAAGATTTGCTCAGGGCTTCGGCGGAGGTTTTGGGGAAAGGGACTTTCGGGACGGCGTACAAAGCGGTGTTGGAGGCGGGCCGGTGGTGCCGTGAAGAGGTTGAAGGATGTGACAATTT[

	G/AJCGAGAAGGAATTTAAGGAGAAGATTGAGGCGGTGGGAGCGATGGATCACGAGAGTTT GGTGCCTCTCAGGGCTTACTATTTTCAGCAGGGATGAGAAGCTCCTTGTCTATGATTATATG CCCATGGGAAGCTTGTCTGCCCTTTTACATGGTCCGTCTTCTTTCTCTTTTTTGTGCTTGCT
>Gm05_ 3828012 4	CACGAGGGCCTAATGTTTCTCATGGAAATATCAAGTCATCAAACATCCTCTTAACCAAATCA TATGATGCCAGAGTGTCTGACTTTGGCCTTGACACCTTGTTGGCCCCCTCTACCCCTA ACCGTGTGGCCGGCTACCGTGCGCCGGAGGTGACTGATCCTCGCAAGGTGTCTCAG[A/C] TGGCAGATGTGTACAGCTTTGGTGTGTTGCTCTTGGAACTTCTGACTGGGAAGGCACCTAC CCATGCTCTCCTGAATGAGGAAGGAGTGGACCTCCCCAGATGGGTTCAATCCGTGGTTAG AGAAGAGTGGACTTCTGAGGTCTTTGATCTTGAGCTCCTTAGGTATCAAAATGTAGAAG
>Gm05_ 3828038 7	GGAGTGGACCTCCCCAGATGGGTTCAATCCGTGGTTAGAGAAGAGTGGACTTCTGAGGTC TTTGATCTTGAGCTCCTTAGGTATCAAAATGTAGAAGAGGAGATGGTTCAGTTGTTGCAACT TGCAGTTGATTGTGCAGCACAATACCTGACAAGCGCCCTTCAATGTCTGAAGTGGT[A/C]A GGAGCATACAAGAGTTGCGAAGGTCTAGCTTGAAAGAGGACCAGGACCAAAATCCAACATG ACAATGATATACTATTATAGAAGAACATGTATCTTCTCCCTCGACTTTGCTTGGCAGATG GGAGCCCATAATTTTACTTGCTAAATCCTGTGCGTATGTTGCTTGCAATTCTGTTGGCT
>Gm05_ 3838682 2	GGAAGCCCACAGGCCTGCAAGCCCACCGTCTATTCCAAGATCTTCAAGACCGCATGCCCC AAGGCCTACTCCTATGCCTATGATGACCCCACCAGCATTGCTACTTGACCAAAGCTAACT ATTTCTCACCCTTCTGCCCCCATCGCCCTGATCCATAATTCATCTTGTATGCTTGT[A/C]T ACTTTATTAGACCATTGGATCTATCATCTATGTATCCCTAAATGGTATTATTATTAATGGT GCTTTGATCAATATTACTACCTTTTTTCGTGAGGTGCGCTAATGTCCTCAGTTTATGCTATA GTGATGGCCAACCACACTCTATTTCTATCCCCACGGTCTAATTGATGTGA
>Gm05_ 3843163 3	ATGCTCTGTGATATATATCATCATGCTCTACTTGATGAGTTCTCAACAGCCCGTGATCTGCT TCTTATGAGCCATTTGCAAGATAGTGTTTCAAGCATATGGACATTTCAACGCAGATACTTTTTA ACAGGGCCATGTCACAGTTAGGTTTATGTGCTTTTCGGGTTGGACTGATTTCTGA[A/T]GCA CATGGCTGTTTGTCTGAACCTACTCTGGTGGAAAGGTGAAGGAGTTGCTGGCACAAGGC GTGTCAAAAGTCGTTATCACGAGAAGACTCCAGAACAGGTAAGTATCCCCCTCCCCCTC TTCAATTTCCGAATGATTGATATGGTGGAAAGTTTATTAGGTGCTGGTCTTAGACTC
>Gm05_ 3843264 7	CATCAATCTAGAGCTCCTGGAGGCAGTACATCTTGTCTGCTATGCTGCTGGAAGTTCCC AATATGGCAGCCAATGTTTCATGATGCCAAACGGAAGGTCATTAGTAAGACTTTCCGTCGCC TGCTTGAGGTAAGTGAGAAACAAACATTTACTGGTCCACCGGAAAATGTCAGAGATC[G/A]T GTTATGGCTGCCACAAGGGTTCTTAACAAGGGAGACTTCCAGAAGGCTTTTGACATTATTG TGTCTCTTGATGTGTGGAATTTGTGAGAAATCGAGACACCGTACTTGAAATGCTGAAGGA CAAAATCAAGGAGGAGGCATTGAGGACATACCTCTTTACCTTCTCTTCATCTTATGA
>Gm05_ 3843274 6	CATTAGTAAGACTTTCCGTGCGCTGCTTGAGGTAAGTGAGAAACAAACATTTACTGGTCCA CCGGAATGTCAGAGATCATGTTATGGCTGCCACAAGGGTTCTTAACAAGGGAGACTTCC AGAAGGCTTTTGACATTATTGTGTCTCTTGATGTGTGGAATTTGTGAGAAATCGAG[G/C]C ACCGTACTTGAAATGCTGAAGGACAAAATCAAGGAGGAGGCATTGAGGACATACCTCTTTA CCTTCTCTTCATCTTATGAATCGTTGAGCCTGGATCAACTCACAAAATTTTTTGATCTCTCTG TTTGTGCACTCATAGCATTGTCAGTAGGATGATGATCAATGAGGAGCTTCATGC
>Gm05_ 3843556 3	TGTCAATTCTTCCCAATTTTTCATGCTCAATGTATATTGCCTTTAACATTGGGATGCAGCTTA GTGTCAAGACTATTAGTTAACAAAGTGTCTTCTGAATGTTTTATAGTTGATCTAATTTCTGA ATATGATTATCAGAATTTTGCAGTGGGGAGAGGACCGACGCTTTGATGAAATG[G/A]GCAGT AATTTGGGTAGACTGGCCATATTTTGATATTTTCAGGTTTCTTAGATTTTAATCATTTTTTTA AGATTTAGCATGCTGATCTTGGTTTTTCTGGATTTATTCTCTTTGATGTCTGTAGTAGAG GATTTACTTCAATTTTTTGCAGGCTGTCTGGGTGTGGGCTGTGAGTTT
>Gm05_ 3843760 9	ACAATAACGAATTGTTTAGGATGGTCAAATTTTGGTGTGTGACTATTACAATATCGAATTGTT TAGGATGGTCAAATTTTGGATGGTCAAATTTTGTGTCTCTGGAATGCTTTAAACTC CTTATTGCATTCCCTTAATTGTAGGTGAAATGGTATTAATTTTTTAGTATCTGC[T/C]CAAGGT GCAGTGCAAACAGTGGGAAAATCAGTGATGCTCTGAAAATTTGGCTAGCACTTGGTTCTTTG AATTTGTTTGGTTCCTGTCTTACGTCAGACTGAAGTTGCAGCTGGTATCATCATCCGCCATG GTTAGGGATGTGGCTTTGTAATGCTATTGCAGCTTTCCTTCTAGATTTAAT
>Gm05_ 3843762 4	TTAGGATGGTCAAATTTTGGTGTGTGACTATTACAATATCGAATTGTTTAGGATGGTCAAAT TGTTTAGGATGGTCAAATTTTGTGTCTCTGGAATGCTTTAAACTCCTTATTGCATTCTT AATTGTAGGTGAAATGGTATTAATTTTTTAGTATCTGCTCAAGGTGCAGTGCA[C/G]ACAGT GGGAAAATCAGTGATGCTCTGAAAATTTGGCTAGCACTTGGTTCTTTGAATTTGTTGGTTCC TGTCTTACGTCAGACTGAAGTTGCAGCTGGTATCATCATCCGCCATGGTTAGGGATGTGGC TTTGAATGCTATTGCAGCTTTCCTTCTAGATTTAATCTTTCTAATTTGCTT

>Gm05_3843772 1	TGCTTTAAACTCCTTATTGCATTCTTAATTGTAGGTGAAATGGTATTAATTTTTTAGTAT CTGCTCAAGGTGCAGTGCAAACAGTGGGAAAAATCAGTGATGCTCTGAAAATTGGCTAGCAC TTGGTTCTTTGAATTTGTTTGGTTCTGTCTTACGTCAGACTGAAGTTGCAGCTG[G/C]TATC ATCATCCGCCATGGTTAGGGATGTGGCTTTGTAATGCTATTGCAGCTTTCTTCTAGATTTA ATCTTTCTAATTTGCTTGTCTCTCTTAGATATGTATAAATTTACGTACTATATATGTTGTGTG CAAGTACTATGGCTTGTGCAGGTAACATAATGGATCAATCATGTTTTTCAT
>Gm05_3858804 1	CCCCACCGTCGTCTCCGACGCCCTCCACAAGGCTGCCGTGAAGGCCGTCGACGTTCTCAC CGCCATGGCCGTCCCCGTGAGCTCTCTGACCGCGACTCCCTCGTGAAGTCCGCTAGCAC ATCCCTCAACAGCAAGGTCGTCAGCCAGTACTCCACGCTCCTGGCTCCCCCTCGCCGTG[G /C]CGCCGTTCTTTCCGTCGTGATGCCGCCAAGCCCGACATGGTCGACCTCCGCGATGTG AAGATCGTGAAGAAGCTCGGCGGCACCGTCGACGACACCGAGCTCGTGAAGGTCTCGTC TTTGACAAGAAGGTCAGCCATGCCGCCGGGGGACCCACCCGCATGGAGAACGCCAAGAT CGC
>Gm05_3858901 9	TGCTGAGGCGCTTGAAGTTATTCCTATACTCTGGCTGAGAATGCTGGTTGAACCCGATT GCCATTGTTACCGAGCTGAGGAATCGTCATGCACAGGGTGAGATAAATGCTGGAATAAATG TGAGGAAGGGTCAGATTACCAACATCTTGGAGGAGAATGTGGTGCAGCCCCCTTCTTG[G/A] TAGCACAAGTGCATCATGTTGGCGACAGAGTGTGTGCGGATGATTTTGAAGATTGATGAT ATTGTAAGTGTGAGGTAGAGTTGATTGAACAATGTGGAATTTAGAGAGAACTGAATCACC CATTATCTTCTTTTTTTTTTTTTTTGTACTTCGTGCTTTAATTTTATGTTGGTAGTTT
>Gm05_3859603 7	GTAGATAAAATAGAGGTAGAGGTGTGGTAGATTTGAGAGCAACATTGAAATGCATGCCAAG ACAGACTCGGAGGTAACAAGCCTGGATGCGTCGTCCAGCACAAGGTCTCCTCGGCGAGCA GTGTACTACGTTTCAGAGCCCTTCCCACGATGGGGAGAAAACGACGACGTCGTTGCACT[T/T ]CACCCCTGTTCTCAGCCCCATGGGTTCCCTCCTCACTCTCACTCCTCCTCCAGCCGCTT CTCCGCTTCACGCCACCGCAATAACCATAATAAATCTTGGAAGGGCATCGACGTCATC GAAGAAGAGGGTCTTCTCCAATCCGAATTAGATCGCCAACATTCTCTCTCTCGTCGATA
>Gm05_3869606 5	AATACCACAGTACAGTGCGTTTCAGGAAAAATGAAAAACCAAGCCCCAGAAGTTCGATTTCAA GTGGGTAACAGAGCAGTAGAGTTGATCCATTACACGTAGGTAGCTACTCTCCTCAGAAGA ACCAATTTTCATAAGCACCATGACGAATATCGAATATTTGAACTTAGTGGAATCAGG[G/T]TC AAACTCATGAGGTGTATTCAAATGGTTGAGGAGGTTGAGGTTCTCATCCACAGCCATGGC TGATGAAGAACTCTGCAATCCTGTTGCTGACCCACCAGCAGCAGAACTGGTAGATGAGGG CGTGTCAGTTAGAGCAAGCACTTCTGGCTCAGTAGGATGCAAGTCTCTGAGAAGCAC
>Gm05_3876518 8	GGAGTTTGACAATGATAACAAATTCCTCACTAAACATAAATCTGCAGATCTGAGCATCCCTAA ACCCACTGAAAAGGCGATTTTTAATTTCTTTCTTGGTTGAGTATGGCAGGTATTGCAATGG AACATTTTCTCTGGACTAAAGAAAACCAAGCATGTGCGACCGAGAGGAATAGCAC[C/A]CTA GGTCTTCTAACTGTTTACCCAGTAAAGTCGGGAAGTTTCCACTGTTGATTTTTCTACCCGG GAAATTCAAGAATGGAAGAAAGAATGCTACTCTGTGTATATCTCTCTGTATTAAGTGTA TTTGTTTATACTTGTTAAAGGGTGCTTTATCAAATCTTGTACCCGGTAACTTT
>Gm05_3886386 7	AGATTCTCTGTCCTGCGGAGCGTCTTCTCTGCGGTACTCGAGCCACACGAGGGGCCAC GTGGACGAGGCACTGCAGCGCGTAGCCGCAGATCCACCAGCGGATCGGCGTGTGGGGT TCTCCTTGACGGTGCAGGCGAGCATCACCGCCGAGACCAGCACGAATGCCATGTTCCACG [T/C]TATGTCGAGCGCCACCACCGGCTTCGAGTAGGTCCAGTCGGCGCGGCGCTCCTCCA GCTCACGCGCCGCGTCTCCCGGACCACGGCGGAGTGGCCGCGGCGGCGGAGGAGGAG CGCTAGGATGGGAGGCGGGCGGCGGCATCGGGGCGCGGGAGAAGGAGCGGCGCGTG GGAAGTGAAG
>Gm05_3889651 4	ATCTCTTGACAGATTGTTCTCTGCAGCGATGTATAGCGCAGTCTCTCCAGCATCGTTCTGAA GAGACATCACATCAGAGAGCGATGACCATTCTTCACTTCTTCACTTTCTCCACCAGTTTCTTC ACCCCATCAAAGTCCCCAGATCCAACGCTCGAAAATATTGATTGGTACGCTAAGAA[C/T]CG CACACCCTTTGCCTCCATCATTCATTCACTTTCTCTCTCAAGTTTCAATTCAACAAACAAA AAACAAAGACAGGGGTTGGTTGTTGTTTTCACTGTTTCAGCTTCCTTCCATCATCAGTTGC AGAAAAGGGTGTCTAAAGTATGAACTTTTTATTTTCTTTGTATCAAAAAA
>Gm05_3908282 3	CCTTTTTTGTAAATTAATTTATTTTTCTTTGAACTTGGTACGGTTAGGGTGGCATGGAAATG TCGAGCAACGAAGTCGGGGAGTAGCCATGACTGCTACAACAGCTCCATCAGTCGAAAATG GCATGAGCCGTTGAATAGTATGCAGTTCTGTAACATAAGACAATGTGTCTACGATGC[T/C]CG TGTTAAACATGAAGAATAATAATGTTTGAAGAATGGTTTCATTCTTTTTTCGATCCCCTTTT CATATTAATAGTTGTCATTTAAATTAATGTTAGACTACTTTTATAATAGATGCGGTATTACAA CGTATCTTTTTCTGTATCCAATTGTGGCAACGAGGATACAAATAAAGTTCTT
>Gm05_	TGGTTCATTCTTTTTTTTCGATCCCCTTTTTCATATTAATAGTTGTCATTTAAATTAATGTTAGA

3908303 8	CTACTTTTATAATAGATGCGGTATTACAACGTATCTTTTTCTGTATCCAATTGTGGCAACGA GGATACAATATAAGTTCTTATGCATTTATCTTAACCTCCGCCGCTAGGCTGAT[G/C]ATAGTG GTTTACAATGCATCTATTTATAAAATAAAATAAAAGACAATGGATCTATTTATCCTTTTCATCC TGTGTGAATATGTTTACAAAATCTAAGGTTTTGTGAAACATGGAATGACTTTCTTGAAAAGT GTGGTTGAAATGTACACTTGACTTATAGAACCAGTAACTCTTGATGTA
>Gm05_ 3908541 6	GATTTCCGCAATTTTCTTTGTAATTGGGGTGAGGTGGTGGTGTGGCTGGGGCTGGGGCTG GGTATACTCTTGCTTCTACCGCTCCTATATGAGACAACAGTACGATTTGCGGGGAAATGCC TGTACGGATTGCTTGATTCATTTCTTCTGCGAGCCCTGTGCCCTCTGTCAAGAATATC[G/A] TGAGCTTCAATTCCGTGGATTTACATGACTATTGGTATCTACCATTGCTTAATTCATTATT ATGGCCCCATAATTTGTTTTGCTTTCCTTTTTCCACATCAAAACAACCTGCTTCCATTACTCT CACTTTCATTTTCTGTTTTGTTACTCCTTCTTCAATTGATACAGCTCATAAA
>Gm07_ 8426860	TATGTATCAGGCTCTTCAATAACTTTACCGGACTCCACACTGGAATCTGCACTAGACTCCAA ACCGGACTCTGCACTAGACTCCACGCCGACTGCACAGAATACAGCCTTTCAGCAGCCTC CCTCATTTCTTTTGGTGGTTCATGAAATGTAAAGCAACCGTAGATAGCAAGACATGC[A/C]T ACAATCCCAAGTGAACATAAACACGTTTAAAGCCTTGATTGAATCAACGTGTAAATTGCGACT AGACTACACTCAAAATTTCCAGACAAGACTCCGCACTGTTTTCTAGTTCCACTTTGTGAGA ACTAACCCCATTTCCATTAACTTTTTTCTTTCTTCAAAAAATAAAACAAAGAA
>Gm08_ 2134000	TGAACCTTCTCCGGATGACCCTTCCATCCTGTGGACAGGAACTCCAGGCTCTCCCCGATCAC CGGGTAACCGGTGGCTCCCGGTGGCAGGTTCCGGGGCCGCAAATGCAGACCTGTGCTTGT AGAAGAGGACGAAGAGGGAGAGGGTCACTATAGAAACGAAGAGGAGAAGGAGGGAGAGA [A/C]GCAAGTTATTATCCTCCATTCTTCTCGTTTTGAGTTAATTTCTTAATTAGTTTTGCTTCGT CACTGATAAATTAAGCTTCTTATAATGTGTTTTAACCCACAGTAGTGAGTGGCATTGGCT TGTGTCTTGGGTGCCAATTTATAAGCACAACTCCTAGCTGGGGAGTGCGCAATGTGAAC
>Gm08_ 2798583	TGTAATCAAGAGGCGTCGGAATCTCAAAGAAAAAATGATGAAACAAATCAAAGTAGCAA TCGGCATTAGATCGGTGAGCTACAGTGACATAAAAAATGAAATAGAATGAACATATGGAAA TGATGAGAGAAAAGAAATAGAAAACCTGAAACACCGAGATCATAGCCAAAGAGAGAAAC[T/A] GCCAAGGGCCCCAACGAAGCATGTATAGACGAAATACCAACTGAACCTGTGCTCGTAAAGA TGAGCCCTTTTTCTGGTCTCCTCAATCTCTACTCCTCCACCAGCCATTCTGTGTTCAAAAG AGAAAAGGGAAAGGTCCACCAAAACACAGCCTTACAGGTTACACATACAGAACTCCAG
>Gm08_ 4388343 7	ACGCAGAACCACCAAGTCAAGCCGGTTCGTGACTCGGATTCTCAATTCCTCGTTCATTCCC GAACGAATTTTAAATTCCGAAGAAAACCGCACCGAACACTGAATTTAGATTCTGAACAAGT TTCTTCCGCGAAACAGCACAGCACTTCAATTTACGTTGGAACAGAGACAAAGGGAT[C/G]C GTGGTTTGAATTTCTCAATCGATTTTCAAATTCGGAACAGCGAACAGTACTTCAATTTACGCT CGAACTAGTCAAAGCGATTCAAATCGATTTTCGCGAACTCGTCCGATATTTCCCTGCAGTG ACTTAGTGATTCTGTTTCATCTTCTCAGCGCGTCTTTCGATTTTTCCGTTAGTCGAT
>Gm08_ 4396066 7	GCCACCTTTGGGTGACCTTTTATAATTTTTTTTTTGTCTACTTTTTGGTGTTTATGTAACACT ATCAATTTGTCATTTTTTCAAAATATTATCCATTAAAAAAGAAATTGTAATCTAATTGTGTAATT GAAACATCTTTGAGGTACTTTCTGTGATTTTGTGAGGTAGAACTATCGGTTTTAA[G/A]GGAGAC ATCAAGGCGTTGTTTCAGTTGCCAAATGTGGGACAAGACTTAAGGATACAGTCCCTGACTGG GAGGTACTACATTTTTTCAATTTGCTTGACGGTAATTTGGTTGAGTCTCAAAAAATTATGATTCAT TTTGGTTGCAAGTTGTGCCACATGGGTTTTGTGTTTACTTTGGTATTGA
>Gm08_ 4396787 4	AATGTTGTTTCAGTACAGCAAAACATGTAAAGCAGCCCTTGATTCTCTTAGTAATGGTGATTT TAAAAAAGAAAAGCTAGTGTGACTGGTACTGGATCATCTAGGAAGCGTTCAAAAGGGTCC TCCAAGTGTAAGAGAGAATGTTAGGGGTCATATCTAGGTTTTGTACGTATGATAAC[A/C]TCA AGCATTAGAACTGACATAGAGACTGAGTGAATTTGTTAGATGTTACACTCAGTCTCCCCT AAATGGTGTCCATGATACTAAAGGCTTTTGTATACCGTAATTACTAATATAGTGGGTATTCTT GGTGTGGCTTTTGAATGTTTCTATTGAAAATTCAGACAACTTGACGTAA
>Gm08_ 4399335 2	TTAGAGCCAATGTTGTTGATTTAACTGCAACAGATATTGGTAATCTTCCCTTGCTTTTTCTTA GGTAGAGGCCTTGGTACACAGGTACATTGATAAGAACCAAAACCTCCACTTAGTAGAAGTTG CAATGCCATTGTTTCATAAATTCTGAGACCCCTTCACTTATGAATACTTGCTTT[A/C]GTAA CATTCCAAGAAGACAAAACAAATTGAGCAATGCAAGTGTGCAAGTAAAGTGAGCATTGGG GATGAGTTTCCAACTCCATAACTTCTTTTTCATATCTTTGGGAAACATTTTCTTCTGCTACC TTTGCAGATATAACAAAGGCTGATTCTGAAAACCCGAAAAACTTCAAGATGA
>Gm08_ 4399426 2	GGACAGGAAATGGAAATAAGCAATTTCAATATAAAATAACCCCAAGTGGTAAAAATGCCCTT ATCTAACAGCTTAAAAAGAGATTGAGTCCCTTAACAAACTATACTTGTGCTGCTAAGTGCTAA GAGGGTCTCATATTAATTCATACCTGTGGAAACAAGGGAATGCCTTTGAGGAGAT[A/C]TAG TGAAGGGATGATGCAGTAATACAGAGTTGGCCAGGAAAGAAGCACCCATAGATTATAGTAA

	CAATATCCCATTGGAGGCCTGGACTGATCAATCCATAAGCATACCATGCAGGACTATACT TAGAAAGCACAATTTGAAAGCCTCCTTCAGACCATCTCTTATGTTGAACTAGTGC
>Gm08_ 4399488 1	AAAAGTTAATTAATGGTCACGTGATCACTTAAAAAATTTACATGCATGTTATCTCTTAATGTT GATTGTTCAAATATGATTTTATAGTACTTATATATATAGTTCTTATTTTTACAATAATGAGCT AAGAGAGGGACAATGAGAAAGAATGTGACCTTTTTTCCCATAATGTGTTTT[A/T]CTCATAG GTACAACTAGCAAGAGCCTTTGATTTTTCTCCAGTTCATGCAAGCTGCCTTCTTTCATATG ATCAATATTCTTATACTCTTCCAATCATTCTTGATTGGTCATTGAACTTTCTTCCGCATAG TATCTCTCTTCTGTGAAAGCAGCCAGTTCCAATATAAAAAGGTCCACC
>Gm08_ 4563265 0	GAAGAACAACAAAAAGCTTCATTTGCCAGCATGAGTTTATTCTTCTGTTCTTGAAAAACAAGG GTCTAAGTGGACAATTAACAGTAAGCAGAGAGGATTACAAAATTTGTCTGGACATAGTGGT TTCTGATTTACAATTGCCTATTTGTTCTGTTCTTGCCACCTCAAATACTACTGTACT[G/A]ATGA CTAGGAAGATTATAGACAAAAAGCATCAATTTGTTTGATTGTAATTTGCAATTTTAACTTG AAAAAGTATTTAGATCTAGATTGTGGCACAATGCATTTTGATCTTTTACTCTTCCCTTTCTTC ATTTAACTTGGGTAGTATTAATACAGATAAAGATCGTTCTTGATGAACAC
>Gm08_ 4569583 5	TCCTTAGATGTTAAGATTGGGATGCATGGATGAACTAAGGATCTGCTCTTCTTGTAATGT CAACTCCACGGGAGGTGTTGTTGAGGATAATTTAGTAATTTAGCCTAATATTCTCGTTCCGT TCTGTTCCAAATGGCATCACTCTCCACCACCGCACCCGGCGTCCCACCCTTCCCT[G/C]CG CCCACCGCCGCCGCGCGCGCTTCAAGACGATTCCAAGCCTCGCCTCGCCTGGACCC GGAACCCGACCCTGAACCCCTTCTCCTTCTTCTTTGTCGGCGTCATTGGTGGTCACCGAC GAGCGGACGCGGAGAATAGCGAGGGAGCTGGAGAAGCTGAAGGCGCGGGAGCGGAAGG
>Gm13_ 5197418	GCATAGCATAATTCACAAAACAAACAAAAAATTAACAAAAAACAAGCGACCAC CCATTTGTCTAAGAGAAGAAAAGGTCAAACCTGGGACTCCACGAGAGCTGAGTGACCCG TGTGGGATCAAACCTCACTGAAGAGCCTCCTCTGTTCAACCTAAGCACCGATCCATG[A/C]C TGAAAAAACCAACACAAACAAATCAAAAACAAACAAAAGGTCAAATTTGCATGCCTTCGCC AACAGAGAGAGAGAGAGAGAAAAAGGGGCAAAAGGGTGTGCGGTATTATCGTCATTACG TGGCCTTGCGCTTGTGTTCCAAACCGGGCAATCGAATTGATGAAACAGAGATCTCGA
>Gm13_ 9272223	TGAATGAATATGCATAAACAATTATTCTCTTGATATTTGAGTAACGATTATTTATTTATTT TGTCAGAACAAGGCACAACCTGGTTGTGATTGCTCATGACGTGGACCCAATTGAGCTGGTG GTCTGGCTTCCAACATTGTGTAGGAAGATGGAAATTCATACTGCATTGTCAAGA[T/A]CTA CTCTAACAATCATGAGCATTGTGCAGCAAGATGTTTGAAAGGGCAGTCAAACAGGAGAATT ATTTCTACGGATGAAGTTAGAAAACACCAGATCTGGCCAAATGTGGACTGTATTAAGGC AGTGTTTATAAAATATCTCAGCATATGAAGTTTCACCTTGAAGGTACTGCTATAA
>Gm13_ 3552426 8	AGTGATCAAATGAATAGTGAACTCTGATTCTGATTGGAGAGTGATACAAAAGTAATTA GAACTTCTGGCTCTAAGTATTGTGCTTTTTTATTTGTGAGGCATTTTGGTTATTGCATTAT GTGTTATGCTGTAATGCATGCAGGCTGCAAGGAATGCTGGCGTGCCAGTAGTGT[C/G]TGA TGCGGGGGGCATGGATGGGCCCTTCCACCACAATTATTGAATTTTGTGATATTCTGAGT CCTAATGAACTGAACCTTGCTCGCCTTACCGGAATGCCAACAGAAAGTTTTGAAGAGATTG CACAGGCTGCTTTGAAATGCCATGAATTGGTCAGTTCAAGTAATGACTTTAGGGG
>Gm13_ 3552436 1	TTATTTGTGAGGCATTTTGGTTATTGCATTATGTGTTATGCTGTAATGCATGCAGGCTGCAA GGAATGCTGGCGTGCCAGTAGTGTGTTGATGCGGGGGGCATGGATGGGCCCTTCCACCA CAATTATTGAATTTTGTGATATTCTGAGTCCTAATGAACTGAACCTGCTCGCCTTA[G/A]C GGAATGCCAACAGAAAGTTTTGAAGAGATTGCACAGGCTGCTTTGAAATGCCATGAATTGG TCAGTTCAAGTAATGACTTTAGGGGTCACAATGAAATGATTCTGCACACTTTTCTACTTGAA TGCCATATTCATTGCTTATGGATAGACAAGGTTTTAGATTGCGGTTATAGTTGCAA
>Gm13_ 3552619 3	ACATAATTAACCTTCATTAGATGCATAATGCATTCCTAGTACAATTCATGAGTCCATTTATTA AAATTTCTCTAGTCTGTTGATCCTGGTCTGCGAACATAGGAGAATCTGTTAGTTCACTTCCC TATTATTGGTTGTTTTACACAGGAAATACAAATCAGTGAAGATTTTGAACGCA[A/C]TAGGA ACAGGAACAGAATAACTTATGCGAGAGAGAAAGACCTTGAAAAATGGCAGCAAGTAAAGC AAAAGCCCTTTTACAGAGCATTTCTGCAAGAAGGATTCTTCAGCATAATAGCTTGTTGTCAT TGTAATAAGATTAGGCCAAGCACTTACCCTTTGGCAATAGAAAGAAAATGAAG
>Gm13_ 3552619 4	CATAATTAACCTTCATTAGATGCATAATGCATTCCTAGTACAATTCATGAGTCCATTTATTA AATTTCTCTAGTCTGTTGATCCTGGTCTGCGAACATAGGAGAATCTGTTAGTTCACTTCCCT ATTATTGGTTGTTTTACACAGGAAATACAAATCAGTGAAGATTTTGAACGCA[A/C]AGGAA CAGGAACAGAATAACTTATGGCAGAGAGAAAGACCTTGAAAAATGGCAGCAAGTAAAGCA AAAGCCCTTTTACAGAGCATTTCTGCAAGAAGGATTCTTCAGCATAATAGCTTGTTGTCATT GTAATAAGATTAGGCCAAGCACTTACCCTTTGGCAATAGAAAGAAAATGAAGA
>Gm13_ 3552619 4	TATATATATTATATATAATAATTTCTTTTAAATTTTGAAGGTCCATTGTTCACTCTAGTACAA

3554978 7	TTCCTCCTAGTTCCACAACCATCTTTTACTTCATACGAGAACAAGCAAATGGCATTAGAGGCCA TCAAAGGTTTTGAGATCAGTAGAATCAAATTCAGAAAGTCAATTAAGGAGCTC[A/C]TGACC TCCGCCAAGAGTGCCAGCAGGTATAAACTATATAATAAATGTGGTGCTAACCTTAACTCTTA ATAAAATTAGAATTTAGAATCAAGAATAGAGTAGGACAACAAAAAGAAACAAAAATGATTA ACACTAGCTCAGGAAAGCTCATAAATCCATATGGTACGTACTATTAGCTGA
>Gm13_ 3558271 7	ATATGATAATATAACTCTACTGATACTTCATTATTAGAAGTGGAACTGATTACATGATGGATT AAGTATTCAGCATACACACTTATCCCCTAACAAATATTGCCTATGGAACAGAACTAACTGTA GTTTTCTAAAGAATTTGAGCATTGGGAAAGCACAAAGGGGAGCTTCAACCTGCATC[A/T]TCAT TTGATTAAACTTGAGCGCTAATATGGGTTTTGATAACATCACACAAGTGCTGAAATTCTGTG CATCATTTTTGTGATCTTTAGCTACATGGCTACTGCTTCATAATCTTCTAATGACAGTTTCAG TGTCATCATTTGTTGAAGGGAATGCTCCTGAAACCTTCAAGGGCTTCAATG
>Gm13_ 3558281 4	TATTGCCTATGGAACAGAACTAACTGTAGTTTCTAAAGAATTTGAGCATTGGGAAAGCACAA GGGGAGCTTCAACCTGCATCATCTTTGATTAACTTGAGCGCTAATATGGGTTTTGATAAC ATCACACAAGTGCTGAAATTCTGTGCATCATTTTTGTGATCTTTAGCTACATGGC[G/A]ACTG CTTCATAATCTTCTAATGACAGTTCAGTGTATCATTTTTGTTGAAGGGAATGCTCCTGAAAC CTTCAAGGGCTTCAATGCTTGCACCCTATTCAATATTTAGATTGATAAGATACTTTAGACTA CTTTCAGGCTCATATGTCAAAAAACATAATATTGATTAATTTCTTTGTGCCT
>Gm13_ 3560010 2	TAATAATGTAAAACTGGCCATCTGTACCTACAACCTACAAGTACCACGTACCAGTACTTGTA CCAGATATATGGTAACTATGATATAGACAATTTCAAGTTATTATTATCACAACTTCTGGTCT CTCTCAAAGATGCAACGGTAATTGCTCTCCTATAAATTGAATCAAACAGACAGG[G/C]ATGT TGCCAACTCCAGTTTCAGACCACATCTAGCAGCAAAAGGTCATTTCTAACTCTGATTCTGTT TTGTGCAGGAACAAGAAGTTGTCAGTGAAGAATAGTCTCAAAGATTAAATGAAGTTTGCCTT TTCTTATTAGCATTACAGATAGACATTGAGATTTTAGATGTAATGTTGATT
>Gm13_ 3561746 4	GTGATAATTGCTGCACGGAGCAACAACGAGATGGATGCATCCAACTGAAGAACGAGTTC CCTGAGCTTCTTTCCATCAAGGAATCACTGATCAAGTATGTCTTTGAGCCAAACAAGAAAAC TGCTTAAATATATTAAGCGGTTTCATGTTTTAGACATCACAGATTATGCATTTTTG[G/A]GG CGGGCTACATAGATCACTCATTATTTTATTGTTATTTTGTGGGTTTTGATAAGAACCATACT ACAAAAGCTAGCCATGTAGTTCTTGTCTCTTAGCTATAAGCGATTCTGTTTCAGCATATT TCCATTATATTACCGTGTGTTAGAGGGTGACAAGTATTATAACTGGGCCAGATA
>Gm13_ 3569074 6	ATTTCTATTGTGCTAGGTGAATTTAACCCAATTTTTATTCTGAATTTTCAGCCTCCTTAATTCT TTATCTATTCAATTTGTAACCTAAACCAAGAATCCTAATTTTTGGTTCCCTTTTGTGAAATAT AGATGGCATGCTATTGCTTCTTGGACTTGGGATGCACAGGACGAAACGTGTG[T/C]GATTTG TAGGATGGCCTTTGATGGATGTTGCTGACTGTAACTTCTGGAGACGACTGCCCGCTG AGTAAGCTACTGATAATTTTACATGGCGATAGACATTGTTATTTTACTTGAATCCATTTGTCT TGGTTAAATTTCTTGATTGATCATTGTGTTTTGGGAATTTTCCAGTAAAG
>Gm13_ 3575947 6	ATTGTCATAGAAAGCAAAGTTTTATTTTGCACGATATAATAGGCGCCAGCTCATTATCCA TAAAGTTACCAGTATTGATATATGCCCAATTTTTGTTCCACTTCGGTGAAATCTTGCTGGA CCTCCAGTATCTGAATCATCATCATATGTGCTTCCATTGTCTGTTACTATCTTTC[C/G]ACCA CAATTTATATGAAGAGAATATAACGCTGTGATAAAAATTAATAAATAAAGTTAGCAATAATTA AAATATTATTTAGAAAAATACATTCCTTCAAATTGTATTTTCTATTCTTTTCTTTCTACTCA AAATAAAGATGCAGTGGAGTTATGTTAACTTATATAGGTCTAACTAGT
>Gm13_ 3578959 0	TTTGCCCACTCTCTTTGTTCTCTCTTCTGTTTAGCCTCTGTTCCGTTTATTCTGCTTGCCTT TTTTTTTTTTTTCTTTTCTTTTTCATTCAATTCTGAAACTCTCCTTTTCAAACCAAGATACG GTGGTTCTTTTCGGATCCTGACAAACCGCCGACCATGATGCGGTTTGTGAT[G/C]CCTTCA AGCAGCATACGTGTGTCGACTATGCTCTTCAATTTTATGTTCTGCTTCCCTACACTTTGG ATCCAACGCTCAACTCATACCACAAGATGAAGGTACTTAATTCATTTCCATTTTGTTTTTT TTTGTTATGATTTTGTATGTTACTATCTCTGTTTCGAAACACTTTTGT
>Gm13_ 3578962 0	TTTAGCCTCTGTTCCGTTTATTCTGCTTGCCTTTTTTTTTTTTTTTTTCTTTTCTTTTTCATTCAAT TCTGAAACTCTCCTTTTCAAACCAAGATACGGTGGTTCTTTTCGGATCCTGACAAACCGCC GGACCATGATGCGGTTTGTGATGCCTTCAAGCAGCATACGTGTGTCGACTAT[G/C]CTCTTC ATTTTTATGTTCTGCTTCCCTACACTTTGGATCCAACGCTCAACTCATACCACAAGATGA AGGTACTTAATTCATTTCCATTTTGTTTTTTTTTTGTATGATTTTGTATGTTACTATCTCT GTTTCGAAACACTTTTGTAACTATATATAAAGTGAAACTGTCTCTTAA
>Gm13_ 3581567 0	GATGTCGGAGGGTTCGGCCACGCGAACTGCGGAGGGCTCATATGCGTATGAATCAAACCG CTGAAACCCTGGGCAAGCCGGTCAGCCAGGTCCTGCCCGTGACGCTGCACCTCTTCCCAT GCCTCAAACGACCTTTCCACAGACATTTTCTTCTTCTTCAACAAAATTCATAACT[G/A] CCCATAAGACCTAGGGTTTCACTTATACTCCCAATCTAATTAGAAGACAACAAATAAAGC

	GTAACGGAAGCTAAATCTAATACTTTTCTATTCTAATCAAAGGAAAGAACAATTCGT CCTATTCAACAAGGGTTTAGGGTTTTCGGTTTTAGTAGTGTTTGCAAACTCAATTTAAA
>Gm13_35819710	ATTCTTATTGGTTACTAGTAATAAAAAACCCTTTACACTATCGACTATCAATCCACGTTCAAATT GATCATGTAATAATGGTGGAGTACCTGGTCTTTGAGTTACCAAGTGTGAGTTCAAGATCATC TGATCCAAATTCCTCATGGATCCTTTCTCCTCCCCAAGGCTTGACAAGCCCCAA[T/T]GTGTT GCTTCCAATGCAAATTCATCTGAAACAGCTTCAGACATAGGAATGTCAGCATTITGATCAG AGCCAGGTGGAATAGCAGGAGAGCATGTTCCACTCTGAGCAGGAGTCCACATTTCGCGAAC CGCTTCCGGGCCAAGGCATCTTCCTTGAAGGCAAAAGGGTTTGAGGAGACCAAG
>Gm13_35823484	AGCCTCGTCCCCGACGTCATCGATCTCAGACCTCAAAAAAAGGTCCTCCTCAA ACCTCCC AAAACACA ACTTTGG ACCAGATTC GATCC ACAA AGTT AAATAG CCTT CTGG GT CGGT CACA AATT GA AAAG GTGT GTGT GTGT AGAAGA AAGAG GTT GTT ATA A[G/T]A AGA ATGT GGAG GATGG TTTG GAAG TGCCA AGTCAGCAC ATTGGT GTGATT TGGGA AGT GGGTTTGTAGCCGAGAGTGATCGGAGATCGGAGGAGATCTGGGTGCGAGAAATGGGAA ATGGTTTTGATGAGAGTGTGTGTGGATGTGTGTGAGAAAGAAGCGGAGGAGAAGAAGAAG
>Gm13_35823512	GACCTCAAAAAAAGTCAAAAGGGTCCTCCTCAAACCTCCCAAAGAACA CACTTTGGACCA GATTCGATCCAACAAGTTAAATAGCCTTCTGGGTGCGTCACAAATTGAAAAGGGTGTGTG TGTGTTAGAAGAAGAAGGTTGTGTTATAAGAAGAATGTGGAGGAGATGGTTGGAAGT[C/A] GCAAAGTCAGCACATTTGGTGGATTTGGGAAGTGGGTTTGTAGCCGAGAGTGATCGGAG ATCGGAGGAGATCTGGGTGCGAGAAATGGGAAGTGGTTTTGATGAGAGTGTGTGTGGATG TGTGTGAGAAAGAAGCGGAGGAGAAGAAGAAGAAAGGAAAAGAGTGAGGAGGAGGAGG C
>Gm13_35823533	GGTCTCCTCAAACCTCCCAAAGAACA CACTTTGG ACCAGATT CGATCCA ACAAGTTAA ATAGCCTTCTGGGTGCGTCACAAATTGAAAAGGGTGTGTGTGTGTTAGAAGAAGAAGGTTG TGTATAAGAAGAATGTGGAGGAGATGGTTGGAAGTGGCAAAGTCAGCACATTTGGT[T/C] GATTTGGGAAGTGGGTTTGTAGCCGAGAGTGATCGGAGATCGGAGGAGATCTGGGTGCG GAGAAATGGGAAATGGTTTTGATGAGAGTGTGTGTGGATGTGTGTGAGAAAGAAGCGGAG GAGAAGAAGAAGAAAAGGAAAAGAGTGAGGAGGAGGAGGCAC TTGATTAGCCAAATTTGC G
>Gm13_35823553	AAGAACACA ACTTTGG ACCAGATT CGATCCA ACAAGTTAAATAGCCTTCTGGGTGCGTCA CAAATTGAAAAGGGTGTGTGTGTGTTAGAAGAAGAAGGTTGTGTTATAAGAAGAATGTGGA GGAGATGGTTGGAAGTGGCAAAGTCAGCACATTTGGTGGATTTGGGAAGTGGGTTTG[C/T] AGCCGGAGAGTGATCGGAGATCGGAGGAGATCTGGGTGCGAGAAATGGGAATGGTTTTG ATGAGAGTGTTGTGGATGTGTGTGAGAAAGAAGCGGAGGAGAAGAAGAAAAGGAAA AGAGTGAGGAGGAGGAGGCAC TTGATTAGCCAAATTTGCGGATCAAATTCATGATCACGG
>Gm13_35835277	TCCTCTTCTCGCCGGAAAAATCAAAAGAAGCTCCGAAAAGAACAAGGAACAAAAAAGAAA CACACACACGCGAACACGAATGTTAGAAAGAAACACAGCACACAGCGCGAGTAAGCGAAT TATAGAGGAAACGACATCCTTTCTGCTTTTTTTATATAAGGTGAGCTAAAACGACGAC[T/C]T CGGATTTGAGAGATTTGGGATTGTGATCTGAGTAGATCTAGGCGAGGGAGGTTGGGTGGT GGAGGCGGAGGAGCATGGTGGCCGGAAAAGCCGCCGAAAAGAGAGGAAGTGGGTTTGT GTTAGGTGTGAACTGTGCACAATGGGGAGAAGAGAAGGTTGAAGGAGCGAAAGAGAAAGA
>Gm13_35849431	TGGTTTCGGTTCCCTCTGCTTATTGGTAAACACAGCGAACAAAGCTCATCTTCTGCTTCTCT TCTCTGATTTTAGGGTAACATCTTTTGTGCTGTCTTTGAAGGTCTTAATCTCTTGCTCTAACT GCTTCAGGTTGTTCTTGAGTTTCGGTTAGGGTTTCGGAGGCAGTTTCTTCTTCTT[G/C]TTCTT CTTCTTCGTCGGAGGGGGTGGGTTTTTGGGAGAGGCCGAGGGCTTGGGTGAAGAGGACG TGGAGGGCTTTTTTGTGGGCCCGCAAGGGTTGGGGTTGGGTTGGGTGAGGGGCGG GCTGGGAGGAGGTGGGTCTGGAAGGGATTTTGAAGGAGGAAAGGGAGAGGGTGCGGCG
>Gm13_35849515	TTTTGTGCTGTCTTTGAAGGTCTTAATCTCTTGCTCTAACTGCTTCAGGTTGTTCTTGAGTTC GGTTAGGGTTTCGGAGGCAGTTTCTTCTTCTTCTTCTTCTTCTTCTTCTCGTCGGAGGGGGTGGGT TTTTGGGAGAGGCCGAGGGCTTGGGTGAAGAGGACGTGGAGGGGCTTTTTGGTGG[G/A]C CCGGCAAGGGTTGGGGTTGGGGTTGGGGTAGGGGCGGGCTGGGAGGAGGTGGGTCTGG AAGGGATTTTGAGGAGGGAAAGGGAGAGGGTGC GGCGAGAGAAGAGGGTACTGAAGATA CCCATGGAGGGCGCAGTGAGAAACCTAGAACGGAAAAGGGAGGCTTTATTTATGGGCTTCT TA
>Gm13_35862124	ACTAATTAATGTCTTATGCTACCAAAGCACTCATACAATCACTAACAATGACGAACACTCAA AATCACTAGTCTTAATTA AAACATCTATCAAATATGTTTCTGAAACTTTGCAATTTCTAATTC CTAATTGCCCAATTTCTTTGCCGCGTTAGCTTCAACOCTAGCCTCGTCATTT[C/T]TTTTGC TTCCGTTGTTACCATTCGGCATTTTTTGAACCACCTCATTATGTTGTCCACCTTTCCCTGG

	CTGATGGGTGACCCTCTTTTCCCTCCAACGCTTCCATGGCTGGGGCCTGACCCTCTTGTTCTCCGTGCGTTCTCACTCTAAACAACAATAACGTCAGTTCTTAATTCATAT
>Gm13_35862205	AACATCTATCAAATATGTTTTCTGAAACTTTGCAATTTCTAATTCCTAATTGCCAATTTCTTTGCCGCGTTAGCTTCAACCCTAGCCTCGTCATTTCTTTTGCTTCCGTTGTTACCATTCGGCATTTTTTGAACCACCTCATTATGTTGTCCACCTTTCCCCTGGCTGATGGGTGAC[C/G]CTCTTTTCCCCTCCAACGCTTCCATGGCTGGGGCCTGACCCTCTTGTTCCCTCCGTGCGTTCTCACTCTAAACAACAATAACGTCAGTTCTTAATTCATATTTTATATATATATGCAGAACATCTAGCTAAGGCTGAATGAGATTTATGCATGTAAGAGTGTTTAACTTGATCACT
>Gm14_27341132	GGTGAACTCATAGATATGAGGCTGTATAGAAGCGGATGCGTGACGTGGAATGGTGATTTGTAGATATGAGGTTGTATTGTTACGTCTTGCTTGCTCGGGAATTTGAAGCCATAGTTTGAAAATCCAACACCTTGAGGACAAGGTGTTTTGATGGGCTCGGGAACCAATGTTACGTCTT[A/C]JAATTTCGAGGACCAAGACAAGCTTATTGAGAGAGAAGAAGAGAGAGCTTTTTTGATATTTCTTTCTTATACATATATATATTGCTAGCTGCTAGGGAACAACTCCCAACAACAATAACAAAAGGATAAGCGCAATCATGGAATATCCTAATGATGATATCCATAACATTCTACACTA
>Gm16_30464934	TCTCTCCCTCCTCATGAGGAGAAAAAAGAAGCTTTCAACTTTTCTATTATTCATCCAACACCATCTCAAAATCGAAAGGAACAAGCTTCACCATCATCATCACCTTTTCACTCCTCATGAGTACCAACCAAAACACCACCTTCTTCCCCCTCCCATGCGCAAAATAACTACTTCTGA[A/C]TTTGTTAATAGCCAGAACAATCCATCCCAGGCTACAAATCAAGGTAACATATATATTATGGCAGGTTGATTATATATCATATTATTCTAATAATTCATATTACCATCGATGTTATTATATGGTCCTCTCAATTATATATATATCATATCATATCAATAATCTGGATCCACATCT
>Gm16_30479527	GCGTGAAAAGCCAGTGTTTCATCAGGCGCCGAACCGAGGATGATATCAAGCTGGCAAACAGTGTTGATGTTGGATCTCTTCGCGATCCCCAGCTGGATGCTGATAGAGTCAAGGACCCAGAA TGGCTCATTGTGATTGGGGTTTGCACCCATTTGGGTTGCATTCCATTGCCAAATGCTG[T/A]TGACTTTGGGGGATGGTTTTGCCCTTGCCATGGATCACATTATGATATTTCAGGCAGAATTA GGAAGGGGCCAGCACCATACAATCTGAGGTTCCCTACTTACACCTTCTTGGAAGAAAACAA GTTGATGATTGGTTGAAAATAAATATGGTTCTGCATACAAGCGGAATGAGTTACAA
>Gm16_30493681	GAACCTTCAGTTCATACCAAAGCCAGGATCGAAGTGCTTACTTCAATCACTCTTCAAAGGA GAGGGATTGTTACCTTATCATCATGATCAGAAAGAAGCTGAGTTAGATTTCCAACTTCAAA CAATGTTATGATGGGGGGTGGCCAATTTCCAGTCATTTCAAGGAGCCATTGCAA[C/G]CGT CGCTGACATTGGACCAGGGGCGAAGGAGAGCTACTGAGGTTTACATGCCAGAAAACATGT CAGAGAATATTTATTCGGATGAAGGAAGGTATTTAATCCCCAGACAAGATCCATTGATCCC CAGGCAAGATCCATTGATCCCCACGCAAGATCCATTGATCCCCAGGCAAGATCCTT
>Gm16_30493887	AGGAGAGCTACTGAGGTTTACATGCCAGAAAACATGTCAGAGAATATTTATTCGGATGAAG GAAGGTATTTAATCCCCAGACAAGATCCATTGATCCCCAGGCAAGATCCATTGATCCCCAC GCAAGATCCATTGATCCCCAGGCAAGATCCTTTGATCCCCAGGCAAGATCCATTGGT[G/A] CCCAGGCAAGATCCTTTGGCTGCAGTAAATATGACTGATTGGGCTGCTAATAATGCTCGCA TAGCAGGGCCTTCCCAATCTCACTTAAATACCTGGAGATTTTATTGGTCATCATTTGGTTCCCT GCTGATCATCAAGTTCGTGGTGGTGGCTGGAATGGATCTGATGGTGGTGGTCTTTCA
>Gm16_30517229	GAGAAATCAGCAAAACCATTTAATCAAACCCCCAAAAAGAAAAGGCAAAAACAATAAGTTG CTACCCAAAAAATATATATGAGAGAAAATCCTCCGAAAAATTCTTTTTTTTTATGAAATTTT GGTTGCATCGACAACATAAGGGGGGGAATGCTCACAGTGGCTGCCGAGACGGC[A/C]GC GTGCGGTGGCTGAGAGTGAGAGGGTGAGCTAAAAGGGGTTGGTGAGGTCTGCGAAACAA AGAGAGGAAATTAACGATCGTACGATGCACCACTAATTAGGAAATTTTCGTTTAAATCGAATC AAATCAAATTGATTTTACAAATCTACTTTAAAAATTTAAATTTTAAATTATAATAAAAAA
>Gm16_30517232	AAATCAGCAAAACCATTTAATCAAACCCCCAAAAAGAAAAGGCAAAAACAATAAGTTGCTAC CAAAAAATATATATGAGAGAAAATCCTCCGAAAAATTCTTTTTTTTTATGAAATTTTGGT TGCATCGACAACATAAGGGGGGGAATGCTCACAGTGGCTGCCGAGACGGCGGC[A/C]TG CGGTGGCTGAGAGTGAGAGGGTGAGCTAAAAGGGGTTGGTGAGGTCTGCGAAACAAAGA GAGGAAATTAACGATCGTACGATGCACCACTAATTAGGAAATTTTCGTTTAAATCGAATCAAA TCAAATTGATTTTACAAATCTACTTTAAAAATTTAAATTTTAAATTATAATAAAAAATT
>Gm16_30539518	CAACAGGCGGACGGGCAAGGTTAACCAGGGTGCTCTTTAGGAAGAAGGGAGATTGAA TGGTTTGTAAACCAGACTTTTCTTTGTCAATGATTGAAAATTTTCGGTTGTTAGGTTTTTGAA ATGGAATCCATTTCCGTTATGGTATATTAATTTTGTAAATTGTAAGACAATATCT[A/C]TCTC AGCCTTCCACAAGACATAGTAGCCAATGTTGAATTTTCTTTATTAAGTGAGTGTGGTTTT GTTCGACAACATTTACTAAAATCAATATGGTTTAGTTTATGTTATTTTACAATAACATGT TTTGTATGCTCTGATGACGTGTTAGTGCAATAATGTACATATGAATTTTG
>Gm16_	AACAGGCGGACGGGCAAGGTTAACCAGGGTGCTCTTTAGGAAGAAGGGAGATTGAAT



3053951 9	GGTTTGTTAACCAAGACTTTTCTTTGTCAATGATTGAAAATTTTCGGTTGTTAGGTTTTTGAAA TGGAATCCATTTCCGTTATGGTATATTAATTTTGTAAATTGTAAGACAATATCTC[A/T]CTCA GCCTTCCACAAGACATAGTAGCCAATGTTGAATTTTTCTTTATTAAAGTGATGTGGTTTTG TTCGACAACATTTACTAAAATCAATATGGTTTAGTTTGATGGTTATTTTTACAATAACATGTTT TGTATGCTCTGATGACGTGTTAGTGTCATAATGTACATATGAATTTTGA
>Gm16_ 3062652 4	TATAATGAGGATGGTGTGTCAGGGAAGAGGTACTATGGTGAAGACTGTGATGTGAGTGATGTA CAGGAAGCAAGGCAATTTAGAGAGATCATTAAAGAGTTGGTGACGTTAGGAGGGGCTAATA ACCCTGGGGACTTCTTGGCTTTGCTTAGGTGGTTTGATTTTGATGGTTTGAGAAGA[G/A]A CTAAAGAGGATTAGTAAGAGAACCGATGCGTTTTTACAAGGACTCATTGATCAGCATCGTA ATGGAAAGCATCGTGCCAATACTATGATAGATCATCTTCTTGCTCAACAGCAATCACAACCT GAATATTACACCGATCAAATCATCAAAGGACTTGCACTGGTAATTAATTAATTATT
>Gm16_ 3067691 4	TCGGTACTTGGACCTCATCTCCTCCATCAGGGCCCGAGTAGTGGGCCCGGTAGCACCGCCC CAGCTGCCTCGTTCGCCGCAGCGCCGCCGCGGAGCACCTCCTCCCGCGTCAGCACCCGCG ACTTCGCCAGCGCCGTGTCGCCCTCGCCGCCCTCGATCGCCACCGGCCACGGCGGAGAG G[G/C]TGCGAACCCTACTCGTCCATCGGTGCCGAGTCGCCAACTCGGAACCGGGTTTCTT GCTGGAACCTCGGAAGAGTTCTGTACGGTAATTTTTATTTTCTGACTTTGATTCTTCGGCGTTG GGAAACATTTAGAACCGTTGGATACCGTTTAAAAATGATCTGACGATCAGTTTGAAAGTAAT
>Gm16_ 3118922 5	ACTCCTTTCTGTTGGTCCCCGACATGGGCGCTACTACTAATAACGGATTTTGAACCATTC CACTCTCATAACTCATTCTTCAAACAAACAAACCTTAAAAACAAACATACTAAACCTTCTC CTTTTCTTTTCTTCTCCATTTTCCCAACTCACACTTCTCATTTTCTCAGGGAAT[G/A]GAAAA AGGAAAATGATCGGAAACTTCATCGACGACATTGACTGCGGCAACTTCTTCGACCACATCG ACGACCTCCTCGAATTCCCCGACGACGCCGCCGCGCTGACACCTCCGCCGCCGCTCCT GTCCCTCCACCGGCGAACTTCTGGTCCGCCGAGTCCGACTCGCTCCCCGCCACCG
>Gm16_ 3119186 3	CAACCCCAATCCAGTGATGCAGCTCATCTCCCCTGCATCCTCCACTGGTGAGAACACGCA GCACAACGCTGCCAACACCTCCAAGGCATCATCGGATTCCGAGAATTTTCGCTGAGTCGGT GATCAAGGCTCCTAAGCAGGCCTCTGGGGAGCACACAAGAAGAAAAAGATCAAAAGTGA[T /C]GTTCCCATCAGGTCAAGAGCGGAATGCACCATCACAGGCAATTAGGAAATGCTTGCACT GTGAGATAACCAAGACACCACAGTGGAGGGCAGGGCCAATGGGGCCGAAAACACTCTGC AATGCTTGTGGCGTGCCTACAAGTCAGGCCGGCTTTTCCCGAATATCGCCCTGCAGCG AG
>Gm16_ 3120445 1	CCCAATCGAAGCCCCACCCATGACTGAGCATGCAATTGCAATTGCAATGCCAGACACACT CTGAAGGATACGCAGGGCATGCCCGGCACCCTCGGTGGCTTCTCTTGCGCTTCGCCAG TTCTCTTTCGCCCTCGTTTCCCTCTCCGTCATGGCTACCACCTCCGATTTCCCTTCTG[C/G] CACTGCCTTCCGGTATTCATTAATTCCCCCCTTTCCTTATTTTCTGTTTAATCGTGCTTATTT TTAATTGGGTTTGTGGTAATTGTTGATTTGGGCTTGTTTTTGTGTTCAAATCAGCACCCC AGATTTTATTTTTTTTCAAGTGGGTTTTATCAAGGGAGCAATGAAATGAACCTT
>Gm16_ 3122568 4	TAACAAAACAGATCATATATCTGCCATTATTCTAAAACAGAAAAGGATAGTTAAAAATAAAGAA TACAAGGTTACTCCAGAATAACGATGTAGTAAAAGGAAAGTATAATGTACAAAGGTTTCTAA ACAATACAATATGGTGAAACAAATAGGATACCTGCATAATTTTCAGGTGCATCAG[G/C]CCCA CACATTGAGGCTCCGAGAACCTTATCTGTCTCAGCATCAACAACAAGCTTCATAACAGTTTT TTCTTGTGCGCTATTTGAAGAATATCAGCAACCATAATCTAGATAAGAAAGTTATGCTGCAA AGAACATAACTAATACTGGGTCCAACTGAGATTGAAGAGCATGTGATGATAA
>Gm16_ 3136345 1	TCGGGTGTTTTGATATGTGCTCGCCCGCCGCGTATCCTATATCGGCTGCGTAACCAATG AGGAACACGGCGACGGCGACGGCTAAGGCGCCCCCGAGAATAAAGGGGCGCCTGCGAC CAAAGCGGGAGGTACAGTGGTGCCTGTAGTAGCCACAAATGGGCTGTACCACGAGCCCG G[G/C]AATGGGCCCGCAGAGCCAGATGAAGGAGGCCGCCGCGTGGGGGACTCCTAAAAG CTGAACGTAGGGGTTAGAAGAGAGAGCTGGAGAGCCCACTGTATGCCGGCGG CGATGGACGCCACGGCAAACATTTTTCGGAGTGGGCTCGCTTGGGCCGGGCCGCCCTCC AGACTAAG
>Gm16_ 3142702 4	AGATTTGGATTGGAATGGTTTCCCCAAGCAAATAAAATTAAATTAATAAATAAAGACAAG CCCTCCTCAAATATCATAAAGCAGTTAACTCACAAGCCTCAACCCCATTTGATCCTACCC AACAAAGAAAATCCCAATGTCATAATTGGTGTCAATTTCAATTTCTAGTGTGTGG[G/A]ATCT TTTTTCTTAGCATGTGTCATCTTTGTTGTTCCCATTTCCATTACCAAGAGCCCTTTGCTTTC TTCATCTCTCTCATTTTCTTATTCTTAAGCCCATTTGCAACTACCCATTTCAATCCTAGTGAG CTTCTCAGCCTTCAAAGCCTCAGCTTCCCAATCTGTCCTCACCACACAA
>Gm16_ 3142730	ATTGCAACTACCCATTTCAATCCTAGTGAGCTTCTCAGCCTTCAAAGCCTCAGCTTCCCAAT CTGTCCTCACCACACAACATAGAGGATTGACACAGCACATGCCACCTGGGCCGACAGAA

4	GCCCCAACCAAAGCCCCTGAACCCACCTTGAACCAAAGGCCAGGCCACGGCCA[T/T] TGGGGTGCCCAAAAGTAGAATGAGCCTAGGTTTATGTGGGCCCAATACCAGGTCGGGC CGTGCCACGAAGGATCCACAGCCCGTGGTTTGTGGACAGTTCCCAAGCTCACACAGGCC CATAATCGGCATGACTGATGCAACCAAGGCTTTTACGGGCTCGTCGTTGGTGAAGGCC
>Gm16_ 3145991 2	TGAGGAATTGTAGATTTGTGAATGAAGAGTTGGGCTGTGGAGCAAGTAAAAGTGCGGGAAT CAACATGGTGTGTATTATGATTGATTGATAACCTTGTCCCTAAGTATGTGGTGAGAGGCCT GATCCCGAGGATTTTGTATTTAGAAAAATATCTGTTGAGAGAGGTGAACCCCTTAAA[G/T]GA ATCTTAGTACTTGAGGGATTAGCCCTGGCTCCGAACGGGAAGATACCTGAGTGCAACCAAA AAAAGAAAAAGGAAGAGTAAAAGTGTGGGAAGAACAAGAGGAGAAAAATAGATTGAG GGCATCTTGGTTTGTAGAACTTAACTGCAAGAATTATGGAGTAATTGATGTTGTTT
>Gm16_ 3145996 9	GAATCAACATGGTGTGTATTATGATTGATTGATAACCTTGTCCCTAAGTATGTGGTGAGAG GCCTGATCCCGAGGATTTTGTATTTAGAAAAATATCTGTTGAGAGAGGTGAACCCCTTAAATG AATCTTAGTACTTGAGGGATTAGCCCTGGCTCCGAACGGGAAGATACCTGAGTGC[C/A]AC CAAAAAAAGAAAAAGGAAGAGTAAAAGTGTGGGAAGAACAAGAGGAGAAAAATAGATT GAGGGCATCTTGGTTTGTAGAACTTAACTGCAAGAATTATGGAGTAATTGATGTTGTTTTG CATGTACTGGAAGAAAAAGAGGTAGTGATTAGATTTAATTTTAAATAGACATAT
>Gm16_ 3146155 4	TGCAGAAAAATGAAGGAAGATGTCATGCTGTTGCAACCTGCTTAACCAGGAAATTTATCTCTT CCACCCTCTACATTTCACTCTTTTCTTGCAACCATCATCATCTCCAAAGTGACCCCAACACC TACTGGTTCAAGTAACAACCTTGATACCTGGTACACCGCGGCTTGACGAAGCCGTG[T/C]C GTGACAAGGGACAGAGTAAGAACTGGAATTTTGATGAAATTGCAATGGAGTCCTAGTCTA AACTATAAGATAGAATAAGACTAGGGAATTTTCTTGTTGGGATAGTTTATAGCATGTTATTGC TATTTTGCCCTTTGAATTTTGTATGGGTGCAATTTATCTCTTCTCTAGTTTTT
>Gm16_ 3147516 3	TATAAGTGTGTTTTCAAATATTAAGGAGTTGTGGGGTGCGAGCGTGATTTTCTCGTTATAA AAATAAGAAGGAAGAGTGATTCAAATCTGAAGCTGAACCCTAACTAGCTGAGCAGACGGAG GAGAAGCTGACTGAAGCTGCCTCTGCGACCGGTTTCGATTACGCTCAGACATTT[C/T]TG TAAGTTCATCGTTTCACACTTTACACAAATACCCTATTCTTCGCATACTTCGTCGCTGCAAC CCGAACAACGATAACCTAATCAATTTGGCTTCGCCCCCTCTTTTCTCTTCACAAACAATA ACAATTCACGTGTGATTGATTTCTCCAGTGTTTTACGAGTTTCAAAGATGCTCG
>Gm16_ 3147635 9	CATTTATCAAAGCCCAGGTTAGCAATGCAAACAGTGAAAGTGTCTGCTTGCTGCTGTAG GTTGCAGTCATGGTGCAAGAAAATGGCAAGAGGAAGAAACCAACATTCTGGTGACTGGC ACACCGGGGACAGGAAAGACAACCTGTGTGCACTGCTCTAGCTGAAGCCACCCAGCTCT[T/ C]TCACATCAATGTCGGAGAATTAGTCAAAGAAAAGAACTTGCAATGATGGATGGGATGATGA GCTTGATTGTTACCTTCTTAATGAAGACTTGGAAGCTATCAATTTATATAATTTGGATCATT CCTATATGCTTGTGTTTGTAGATTTAGCTCCTCTAAAGTAAAATGAGGAGATGTGTAA
>Gm16_ 3147995 8	TAGGACTTGTAGTTGGGGACACCATGAGACTATTAACCCTTCTCTGATGTGTGTCAGCAAC TCCTTTGGAAGCTTTCCCTTATCACAATCTAATCACCCACATGAAATAACTTCCACTATCT CCCAAACCCCAAGCCAGTTTCCCTCAGTTTGTCTCCTCATTAAAGCCCTGCCATGCTCC[G/C]GAA AGAGACATACACAACCTGACCCTTTTGGTTTTTTCATCAAGCCATTTGATGCAAGCTTCTGAGT TTGGATTATACATGTTAACACCATAGTCCTTGTCTCTTGGAGTCTTTTGTCCAAGTAGATA GATGGCAGGCATGGTCCTATTGGCTTTAATGGCCAAATCTTCACCAGCCAATC
>Gm16_ 3150633 3	TAAGTAACGATAACAGGGGCCACAAGAGAGACGTTAACTAGCAGCTTCATTTGTGCTACA AAATATTTGAGGTCACCAATTGAGGACACTAGAACCATAACAAATTTCAAATGCGCATACAA AATTTTCTTCCACTCTTCAAACACGCGAATTCTAATCTGAACCTCAACACTCCG[G/A]GGA ACAAAAATGCTACTCCTTCCCGTGCTTTCTCTCTCTCCCTCCTTCCCCTCTCTCGCTTCGAA CCGTTATCTCCAATTTGACCTCGTCGGAGCCGCCGCGTCGTCGATGCGCCGCCCGGGA GTCCGGTGGCGATGAGTTCACGGCGAAGTCGGGGTACCTGTTGAGCTGAGCGCC
>Gm16_ 3150658 7	CCAATTTGACCTCGTCGGAGCCGCCGCGTCGTCGATGCGCCGCCCGGAGTCCGGTG GCGATGAGTTCACGGCGAAGTCGGGGTACCTGTTGAGCTGAGCGCCACGGAGGCGGAT TCGCTGGGGGAATACCGGATACCGAAAAATCGCGGCCGTATACTCGCGGAAGCCTCTGCTC G[C/T]TGCTCGCCGCTTGGTGACACCGGCTCGCGTTCGGGAAGTGGTTCGGTCTCCGC TACATCGATACTCTTGGACCGCTCCGAGTCAATGTTTCAGGTTTCGTAACGCTCTGTTTG TTTCTATTTGTTATTTATTTTGGCGAATTCAGGTTTGGAGGAAGTAGTTTCGTGTTATTAGA
>Gm16_ 3151605 2	TCGTAAATTGCAATATGGGGATCTTTTTGGATTTTACTTGTTTTATTATACCTGACTTTTAG TCAATCGTTCAGCAATAGGTATATTAACCTTGAACGCTACATGGACAATCATTTTACTGTT TTGTTTATCCTAATAAAAAAATTGTACTTTTTCATCAATGACATGTAGGAAAGC[C/G]GACAG GCGAGTCCTGACAAGAATCAAATACTGTACATGGAGGAGTTGTCGTTAGTTTTAAATCAAG TTGAATCCATTCAAGACATTCTTCTATAATTTCTGTCATTCTTGAGGTAATCTAAGCATTA

	GCATTTTATTTAAGAAATGTGGCTGTTTACCCATTTTTTCATTAACCTGTA
>Gm16_31516059	TTGCAATATGGGGATCTTTTTGGATTTTACTTGTTTTATTATACCTGACTTTTAGTCAATCG TTCAGCAATAGGTATATTAACCTTGAACGCTACATGGACAATCATTTTACTGTTTTGTTTAT CCTAATAAAAAAATTGTACTTTTCATCAATGACATGTAGGAAAGCAGACAGG[C/T]GAGTC CTGACAAGAATCAAATACTGTACATGGAGGAGTTGTCGTTAGTTTTAAATCAAGTTGAATCC ATTCAAGACATTCTTCTATAATTTCTGTCATTCTTGAGGTAAATCTAAGCATTAGCATTTTA TTAAGAAATGTGGCTGTTTACCCATTTTTTCATTAACCTGTATAAGTCA
>Gm16_31520234	TTGATCATCAATGGGTTTCAATTGAACGGAAGATCTCACTTCTCAGTTGCTGTTTCATGCAT GTATATTTTTGGATAAAATTCAAATTTGAAGATGTGGATCTAGTGTGAGAAAATTAATGAAT TGTGTGCTGTTTTCAACCGGTGTCAATAACATGGACCAAATGGGAATGTTAGCC[G/T]TAGA TTGAAGTCCATGTTTTGTAACAAAAGCTTGCAGGGAAGTATCATTTAGCCTAACCTTGTA AGGGGCAAACTAAATTATGGATCTGATGAATTGGTGAGCAAGTGAAGGAAAAAATATATG CAGTGGGAACAAAAATTGCGTACGGGACAAAAGGCACCAAGCATCATTGTTTA
>Gm16_31520242	CAATGGGTTTCAATTGAACGGAAGATCTCACTTCTCAGTTGCTGTTTCATGCATGTATATTT TTGGATAAAATTCAAATTTGAAGATGTGGATCTAGTGTGAGAAAATTAATGAATTGTGTGC TGTTTTCAACCGGTGTCAATAACATGGACCAAATGGGAATGTTAGCCATAGATTG[T/G]AGT CCATGTTTTGTAACAAAAGCTTGCAGGGAAGTATCATTTAGCCTAACCTTGTAAGGGGC AACTAAATTATGGATCTGATGAATTGGTGAGCAAGTGAAGGAAAAAATATATGCAGTGG GAACAAAAATTGCGTACGGGACAAAAGGCACCAAGCATCATTGTTTACTTCTCTA
>Gm16_31520380	GTCAATAACATGGACCAAATGGGAATGTTAGCCATAGATTGAAGTCCATGTTTTGTAACAAA AGCTTGCAGGGAAGTATCATTTAGCCTAACCTTGTAAGGGGCAAACTAAATTATGGATC TGATGAATTGGTGAGCAAGTGAAGGAAAAAATATATGCAGTGGGAACAAAAATTG[G/A]GT ACGGGACAAAAGGCACCAAGCATCATTGTTTACTTCTCTAGCATTGAAGTCATTGATCAATG TGAAATGGGATTAGAATTGATGGCATCAATATGGTTAAGTAGCTGTCCGTGTTCAAGATAA CTACGCTTGCATGTGCGCATGTATTTCCAGCTTTCACATGATTTTCATTCAAATA
>Gm16_31525076	TTGTTTTCCAATGTGAAAGAACTATACCTAAATGGGAATGATTTACAAATCTTCTCCTGCCTG CATTCAAGAATTTTCAGTTTCTGACAGAACTTTATTTGGAAGCTTGTGAGAATCTTCATGAAAT TGGATGGATTCCACCGAATTTAGAAATATTCGCTGTAATAGATTGCACATCCTT[A/G]AAAGA ATTAGACCTCACACTTCTTCCACATGGAACAAAGAATGTTGCCTTTTAAGGAAGCTTCTTT TATGTGGATGCCGAATCTTGAAAACTTAAAGGGATTCCACTGAACATAGAAGAATTGATT GTAGAATCCTGCAATTCCTTGAAAGATCTAGATCTTACTCTTCTCCTCTCA
>Gm16_31525931	ATTATTGATCACATAGAGGAACTATTCAATGACTGTGTTCTTTTCGAAAAATGAATGGAATCA TGTGGTGTGTACTACATCATGGGTACCCCAACCAATCAAACAAATTGGAATCCATGTAATCA AACAAGGAAGCAACTTGGAGGATATCCAATTTACCAATCCACTATTGTCCAAAGA[T/C]GAC CCAGATTTTCGCATACAACATGATGATTCAAGGAAACGAAAAAGACAAGAGTGACTGTCTT CTCCCGGAGATGAAGTATCTTCCCGTTGATTGATAATAAAATTTGTGGTTTTAGGACACCT TGTTTTGATTATATATTGTGCAATGTTATGTATTCCCAATATAATTACAGGAGG
>Gm16_31675129	ATTAACACGTGGCACAGCAAGATTTGAGTCCAGATTGTTAACCATTGTAGCCACTTTGTTAG CATCACCATCCAATTGACAATAATTTTGTATCAACTAGTACATTCATCTTTTTCAGTGT TGATAAGAAAGTTTAGAATCACTAGGTATTGAGTAATGCTGTTTGATCATATG[G/A]AAGGT GACATTCCTCAAATGTCAATATATTCCTGAAGAACACTTCTGAATTGTCAGTTATATTCAAGA TTGGCATTGTCAACACACCCTCATCGGAATAGGTCATGTCAAGTAAGCATTCAATTTTCAAT GGACTGACCTTGAATTTAAGACCTGCCTCCCGTAGTTGGCTTGCACTGTA
>Gm16_31675346	AACACTTCTGAATTGTCAGTTATATTCAAGATTGGCATTGTCAACACACCCTCATCGGAATA GGTCATGTCAAGTAAGCATTCAATTTTCAATTTGGACTGACCTTGAATTTAAGACCTGCCTCCC GTAGTTGGCTTGCACTGTATACATGTTTTATACCTTTGCATTCTTTTTGTTCTCT[C/T]GGAAC AAATTTTGATGACGATATTATACTAGTTCTCATAAGATCGGTGAAGTGCTTTGGGGACTCTG TTGGACACGATGTTGTTCCATATCCACATGCTTCAAACAGTTGAAAGAAATCTGAAGAAAG GTGGGAAATTCCTGATTCATACCAGCGATGTTGTAAAGTTGTTCAAGAACA
>Gm16_31708589	AATTGACAAATATCTTTTATATTTTACCTTAATAAGAAGAAAAGGACATAGGAAGG AATCAACACAATGCATGTTATGCCACAATTGCTAGAAAACGATACAAAGAAATCTGGCAGTA GAAAAAACAAAACTCCCTGTCAACCCCTTAAATCCCTTACATACCAATTACAT[T/A]TATCA CTCCAAAATTCCTACACCATGCCTTCACACAAAACTCCCTCGTGCCTATATATGAAACAAAG CCATTGATGCAAAAAATATGAAAAGAAGAAAAGGCAAAGGGAAAAATATCTTACATACAATGC ACACCTCAATCACATAATGTACATATACAGGATATAGCCAATCAAACAAAC
>Gm16_3170966	ATGCATGCTGACAAGTTTATACACGATATAAGAAGCAATTCACCTGTCCAGTTAAGCAGCTA AATTCCAATTACTTAAATTTGTCATGACTTGTTTCATTTTCAATCTCTGCATCAGTGAAAGTTA

3	GGGCTCGCTGTAGCTTCCTAGGAGAAAAATGTCTCAACTTCAACCATATATTTGG[A/C]AGTC ACAGGTCTGTGATCAGACAGTTTGAGCTCTGCCCTTCTGTAACCTCAATAATCTCATTCCCTT GCCATATGAAAGAATACGATCACACCTTCAAGGTGATTACAAACAAGCAGAAAAATTCATCAA GGGTAAGACATGTATCAAATACCATACCCTAGTTCATAAACCATGAAATGG
>Gm16_ 3173192 4	TCTCAACGACCAGATGGCCTCCATAACAATGTCTCCCTTTTGAATGAAGCACTTTATGC TAACCTGGAATGGTTCAAGTTTGAGAGATAATGATGAAACAATTACCCTTGAAAGCATAATA ACATTATTCATAGGAAAGCTTGATTGATAGAACCTGACATTGCATAGCACAAAAA[G/T]TTCC AAAACGGGGGATTTCATTCTTGGAAGAAAACATTGAGTCTATAACATCCAAGCTTTTGATTG AGTCTTTATTCATTATAAGTTGATTGCATGCATTGCAGAGCACACAAGGTATGTGGTTTCA ATAGTGTGATATGCTTTCAGATTATTTTTTTCCTTTCATTTTTGCCTTAA
>Gm16_ 3178995 8	TTAGGTATAGTAAAAAAATTTGGCTCGTGCAAGAAGAATATTCCTTTTGGAACACGAGCG AACAAGCAAGCTCCTTCTTCCCGTTGAGTTAAAGCAATTATAACGTGGCTCTCATGCTTT CCATACTTACCCTGCATCCCTCTCCCTCTTCTCCTTTTCTCCATAATTATAAAA[G/T]TCGC ACCCAAACACCCCCCTTCCCTCTGCCACCCAACCAATCTCGATCTCGCCGAACAACCCCA CCCTTTCAAATCCGATTTTCCGATGCCGAAGAAGCGAAAGTCTATCGCCACCAGCCTCGA CGAGGTGGATCGAACCCCTCTACGCGTCGTTTTGCACCGCCGCCAACTCCCTCTC
>Gm16_ 3179007 8	CTTCCATACTTACCCTGCATCCCTCTCCCTCTTCTCCTTTTCTCCATAATTATAAAAATCG CACCCAAACACCCCCCTTCCCTCTGCCACCCAACCAATCTCGATCTCGCCGAACAACCCC ACCCTTTCAAATCCGATTTTCCGATGCCGAAGAAGCGAAAGTCTATCGCCACCA[A/G]CCT CGACGAGGTGGATCGAACCCCTCTACGCGTCGTTTTGCACCGCCGCCAACTCCCTCTCGCA GCTCTACACGCACTCCATGAACCACCAGAAGCTCTCCTTCAACGCCGGGGAACGACACGC CCTCGTAATATAATAAACCTGCCCTCCTTTTTATTTCTTTCTTTGTTATTCTC
>Gm16_ 3179187 3	AATGTTGTAAATGTAGTGTCAAATGTTCCATTGTGATGCTTAATTCTTTAAATTTATTGATA TTTTCTTTGTTTTGGTTGTAAGTATGTGAAAGTTCAATCCTCTTGCAAGATGAGCTGGATTAT TGTGGAGAAGAACCCTATGTACCTAGACACCACTGCAACACCACACAC[A/G]GCAG CATTGCATGTCCCTAGTTTTCCAGTCACTTGCAGCATCTTCTGGTCAAACAATAGCTGCACAA GGACTCCGCTCGGACCACTGTGAAAATCAATCAAAGAATTCTGTGTTTTCAAATGCTTTGTC AAGTCCAGTACGTCGAAGCCTTCAACATTATCAAATTGGTGAGGGAGGATGC
>Gm16_ 3182788 4	GAAACAACAATCCCCAAAATAGAACACACCCATATCCTTAGCAATTAACAACACCACACAA GAAGAACTACACATAACTTACAACTTTTGTCCATTCAAACTGATCAACCAATTGCACCA CCTCTTCCACCACAGACGTCAACAAAGACTTCCCTAGCATTACCACATTGCATT[G/T]CCA CGAATTTACCTTGATTTCCAGTGAGCACATCCAATTGACTCAACTTTATAGTAGCATCCAC AACTTCTCAAAGAACAACGTTTGGTTCAACGCAAATGCATTGACCAACCCCTTAGTCCTCT TATCATTGAGCAAGTCTGGTCAGAGGTGAACACCCCTTGCGGTTTCATTAGG
>Gm16_ 3182799 1	ATCAACCAATTGCACCACCTTCCACCACAGACGTCAACAAAGACTTCTAGCATTACC ACATTGCATTGCCACGAATTTACCTTGATTTCCAGTGAGCACATCCAATTGACTCAACTT TATAGTACATCCACAAACTTCTCAAAGAACAACGTTTGGTTCAACGCAAATGCAT[T/A]GAC CAACCCCTTAGTCCTCTTATCATTGAGCAAGTCTGGTCAGAGGTGAACACCCCTTGCGCG TTCATTAGGTCAAGGTAGTATTTGTTGTGAACACAGTCGGGGTTCTGATGTCCAAGTTGA CGGTGTTGCCGGAGTTGCGTCGGGGCAAGTGGATTGTAGTTGCTTGGCTAGGGT
>Gm16_ 3182813 7	CAAAGAACAACGTTTGGTTCAACGCAAATGCATTGACCAACCCCTTAGTCCTCTTATCATTG AGCAAGTCCTGGTCAGAGGTGAACACCCCTTGCGGTTTCATTAGGTCAAGGTAGTATTTGT TGTCGAACACAGTCGGGGTTCTGATGTCCAAGTTGACGGTGTGCGCGAGTTTGCG[A/G]C GGGGCAAGTGATTGTAGTTGCTTGGCTAGGGTTTTGTCCATGTTGGGGTCGAGAGGGGA GAGCCTGTTGAAGAATGTGCCGCAATGGGCACGACCAAGGTGTGTGCGCCGGATAAGG CGACAACGTCGGTGACATCAAAGTTTTTGGCGCGAAAGCGTCAAGGGTCACGCCGGTGG
>Gm16_ 3182973 8	ATATCACAAGAAAAGCTTCAAACTATAGTTAGTAAGAGCAGAGGGTTTTGAGAACCAAAA TAGTAGTCAAGTAAAGGTAAAGGGATGAAGAGATTACAAGGAAAATCATAACCGTGTCT GCCAAAACAGTGATATCTGCGCAAGAAACGATCCTTCCACACTCCTTATGGATAAT[G/A]GC TCTGATGTCAATGGTTTGCAAGGCCTCAGTTCTGATGCCTCCGTTGGCTGGTTGGTCT CTTTCATTGGACTCCCATCTAGCAGCAATGATCCATCACATCCCTATACAAATGCAAATTA AGTTAGAGGTAATATCTTTTTCTATATATATATATGCATGCATATATATTAACC
>Gm16_ 3184075 3	GTTGGGGGACTCGGCATTGGGACACGTAGCTATCAAATTGTTGTTGAAGTTTGGGTGCGATT GGTGGGTCCGTTTCGATGGTTCTGTTCACTAGGGAGGGACAGTGGGCACGACCGTAGGTG TGTGCGCCGGAGAGAGCGACCACATCGGTGGCATCGAAACCTCTATTTCCGAATCCTC[C/T] JCAGAAGATCGTCGGTTCGGAAGAATGGTGCCGGTAGGTTGTCCGGCGCGGTGGCGTTTG GTCCTAGGCCGTCTTTCTTCCAGTGGCACGTCAAATCAGGGCCTCCTAACTGTTCAAA

	TTACAAAGAAGTTACCAAAAGGAACAAACGAGAGATCAAGCATTATGTTTAAATGTGTTT
>Gm16_ 3184081 9	GTCCGTTTCGATGGTTCTGTTCAGTGGGAGGGACAGTGGGCACGACCGTAGGTGTGTGC GCCGGAGAGAGCGACCACATCGGTGGCATCGAAACCTCTATTTCCGAATCCTCTCAGAAG ATCGTCGGTTCGGAAGAATGGTGCCGGTAGGTTGTCCGGCGCGGTGGCGTTTGGTCCTA[ C/T]GCCGTCTTTTCTTCCCAGTGGCACGTCAAAATCAGGGCCTCCTAACTGTTCAAATTACA AAGAAGTTACCAAAAGGAACAAACGAGAGATCAAGCATTATGTTTAAATGTGTTTTAATCT CATTATAATGTTATAATTTGAGTGAAAATTATCTCTAATTTATATATTATAAATGATTT
>Gm16_ 3184281 5	GAGCTACTCCACTGTCCTTCTCGAAGACATCCTCAAGATGCTTCCTTATAATCCTCTCAAGC TTGGGACATGTCAACAAATAGTAATTCCATGATAGTCCAGGGACTAGTTTCCCAGAGGAAA CATGGATTTGAGAAGCCAATACAAGGGAAGAAATCAAGAGCAAAGAGTGAAACAAA[T/C]G ACTAACCCCTAGCCATGTTTGATTAATTTGCTATGAGTGTGACTTGAGATAACCAAACTTAG AAGAGTTATTTATAGATTCAATATTGAGGTAACGTTCTTCGTTGCTTCCCCTTCCCACCC ACTAGTCTTCTACGTATAATATTTAAACGTGGCGCAAGCGTTTTTTGAGATTTG
>Gm16_ 3188512 3	TGTCTTGGCCGATTTTCTCTCTCACTCTGTACACAGAGACAACCATGCCGTGGCTCCAGGT CTTCTTATTTTCTTAACATTATGTTATTTCCAGATGCCTCAAAAAATGGTGTGCACAAGAGAC AATTCAAAATACTATCCGCAGAGCTAGCAAGGTGATGACGGAGCAAGAGGCTCGA[A/C]GG ATTCTTGGTGTTACAGAGGAAACTCCCTGGGAGGAGATTATCAAGGTTAGTATATCATCTTT GTTGTGGGTATATTACCCATTTGTTTTCAATTTAGCACTTCTAAAGTGGGGAATGCATTTT GAGGATAAAGCACACTGTTATAGCCTTCAATTGGAAAATCAATAGCTAACATAT
>Gm17_ 7914641	TTAAATTTTTTTCATTTGCATGATTTCTGTTTCAATTACTCTATCATTGTCTTGTCTCGTCAT TTTACATAAGGTTATATTTGCTTATCCGAATTAGCTGAGGGGTTAAGCTGGTTTTTGAATTTT GTTATCAGTACACCATTGTCTTTGAAGTGATCATCTCTCTTAAACCATCAGA[A/C]TGACTTG CAGGTATATTTTAGGATGTTAGATGCGTTGTTGGCTGAAGAGAGAATTTCTGATGAGATTTT TAGTCAAACCTCAGGTTTGTCAACTATATTTCTTATTTCTGATGCGTTGTTGTATGTTTTTCA AAATGCATACTCTGATGGATCCTGTAATAGTTGCAGGTTTTATTATGT
>Gm18_ 2151094	ATTGGCGAAATGGAGGACAGACCGATGATGACTGGGTTACATACCGTGGCTGACATTTTCT GTGTGGGTTGTGGATCAATCGTGGGTTGAAATATGTAAGATTATGAGTCAGTGCTTCAAA CATCATGGTTTTTGCCTGTGTTTATCTAGTCTTCAGTTACTCAAGTGGTGCTTTTTC[A/C]GG AGACTGCCTATGAAAAAAACCAGAAGTACAAGGAAGGAAAATCAGTGCTTGAACGGTCAGT AGAAATGCTGTGAAAAAAATTGATACTAACATAATCAAAGAGAAGTCATGTTTCAGTGTTTAG TATGTAGTGTTTAGATAAGTTCTATGATTGGGCAGGTACAAAGTGTCAAGGTCCT
>Gm18_ 3731008 4	TTTATTAATTTAAATTCAAAAAGATAGATGAATTGAGCAAAAGCAAAAAGATTCAACAATT TATAAGAATAATCCAATGATTTCTGAAACAACCATTTCAATTCAATTGATAAATTCAGTAAAT TGTTTCATTGAAAAAAATAAAATAAAGGATCTTAATGCTAAAAGAAAAGCAGTT[T/A]TAAAA AATTGAAAAAATTAATAAGGGGCTTGTAATTTTCAAGAGACAGATATTCATTCAAACAAAACAA CTTATGATAGTAAAGGGTAGAGTTCGAAATTTTTTTTATAGATATTATAAAGAAGAAGAAAT GCTCGATTAACTCGAAGATCAAAATCTTTTTTTTAAATTTTTTCATGAAA
>Gm19_ 6362210	CCTTCTTCTCTTCCCTCCCAACCATCAGATACCCAACCTGGCTGCTTCATATTTCTCCACCTCA GCTATATATATATATATATATATATATATATATATATATATATATATATATATATATATAT ATATATAAATCAATTGAACCAAAACCAATCTCACAATAAAAATCACAATCCT[A/C]ACCTTCAA CTTTTCCCTCAGTAAATCAGCCTCCTGCAATTTCTCCTTTATTTTATTCTCACAATCTGATAT TGCATTTTTCGAACGCAGCATTCTCTTCTTCAACGTCACCTCTCTAAGTTGGGCCTGCACC ACAATTGCATTTAAGTGTCCCAAGTTCCAATTGTATGAATTGAATCCC
>Gm20_ 2831907	TTAGTACATAGCACTCTACTTGTCTATGACATGATGATGAACACAAAAGATATTACACGACA TGCTGGAAAATCTTGGTTTTGAAATATAATTCTCTTTATTTCTGTTATTTGACTGCTAGGAGA CTATATTGCCTGCTAATACTATGAGTGCAGTTGACTTGTGAACTTTTACTCTTT[A/C]AGCGA CTGCATTGTGAATTAGCTGATGCCATTATACGCCAAATACCCGATTGTGGTCACCTTCCTCA TGTTGAAAGACCAGATTCTGTTGTCAAATTCATTGTTGAATTTGTTTCAGAGCAAAAACCAACA CAGAAAAGGAATGTGTTTACCCCGTAGACAAGGAAGCATGCTCTGTCCAT
>Gm20_ 4630295 7	AGACATTATCTGGGTCAATTCTTCTGATTATTTTTGAGCTTTGTAAACTCCAAGTTGTCTT TGAACCTCTCAACGTCCTCATTCAAACCCCTCCAACCGATTCCAATATTCAACGCGGCTCTT TGTTCTCCTTGGACAGCCTCCACAATCCTCCTCTCCGCATAAGTAGTGAATCCC[T/A]AAAA CCAAAATCAAGAACAACAATCTCATGATCGCCATCAAATCAAACGCACTAGTAGGAGAAT ATTACCTCATTCAACCAGAAATGTTTCATTGGTCTTGTTAGTGATCAAATCCCAGTCCAAC GTGAGCAAGCTCGTGAGCCACAACCTGTGCACCGGTGGCATCCCCCTTAATAA

**Supplemental Table S4.** SNPs identified by Method 2 on the four-library comparison

This table contains 469 SNPs identified using Method 2 on the four-library RNA-Seq comparison between the Clark/IsoClark 19-day root and leaf experimental data. The file contains a unique SNP identifier that includes the genomic location. The sequence surrounding each SNP based on the soybean Glyma1.01 genome assembly is also provided.

SNP	[Clark/IsoClark]
>Gm01_1 840408	ACCTGTGCCAAACCTGATTGCATTCCAAATATCTGGCTCTTTCTCCCGTGAGAGGTCAATGC ATTTGGCATAGCAACCACCTTCAATGTTGGAGACACCATTGTCACCCCAACAGTGTTTCATCA TCTCCGATTAGATACCTATTGTGATCGGTCGAAAGAGTGGTCTTTCCGGTACCTA[T/C]AAC AAACAATAAAAAAGAGAACAAAAGGATCCATTAGGATTGAATATAAATGGAGACACAAGTAG CCATAACACAAAACATCCATGTGTAAGCATGCAACATAGTTGAGATAATTAACAAGCAGTGG AGTATTGAGTATGGCTGAGTAATAAGAGGGGAAGGAACATATGGTAGGGTAGG
>Gm01_4 6870875	TGTGCTTTGCATAAGGCAGAAAGTATATTAGGTATAAGCCTTTTATGGCTGAGGTTAAGGACT ATGCTTGCCATAGGGCTGGGGAGCCTGCTGCTAAGAAATCTAAGGTTTTGAAGAGGGAGC CCTATGCTTCCAAGAGGTGTGGATGTGGGTTTAGGATTAGGGCAATTGTGCCGATTG[T/C]A AATTATAATGAGAAGGATAAGAGTTTTGTGTATCAGGAGGAAGGGGTGGCGGTTTTCAAGC TGTATGCTGTGCACTCCGGGCACGAGCCGGGCGGTTGGACGGGAATGCAAGGATTATGC ATAGGGTTGTTGGGCATAAGGGGGGTGGGTTTTTATGGATCAGGAGAATAATGTTTA
>Gm01_5 3218289	GGTCTTAACATGGAGAGGGAAGTAAAAATTAAGTGAGAAATAATTTAGCTTCTCCTTAT TTTTTCTTCTCTCAAGTTACCGGTTCTAGTCATGAGTTGGACCAATAAGGCCCCGCCATT TACTTTGGATATATCTAAGCCCGCAACCTTACCGTAAAACCCTAGCAATACTAA[G/C]AGCA AAAACCTATTTCGCGATCTCGTTCTGTACGTGCAGCCGCAGCAAGTGATCGAAGCGAAAAGA GAGAAATTCATCGTCGCAGTAATCATGGGTAAGTTGATTGAAAATTAGGTATGGTAGTGT CCCTTTCTTGCGCACGACGAATAACAACATATAGAACGCAATTTGCAGGTAAG
>Gm01_5 3669556	AAAGGAAACGCAACTGGCACAGTAACCTCTCAATCGGTTACCTCTTTGAAGTCGAAGCGCAA GAGGTAGACATATTATTTTGTCCCTAACTATGTGGAACAGCTAGCAAGCAACACAACATAT ATAAGGACATGCTTAATATTGCTGATAGAGTGTAGTCCCACTAAGGATGCAAAAGA[T/G]GG TAGATGCATGTGGGTTACTGTAATATTGTAAGATAGCTGAAACAAATTATAGACTGGCTGTA AATTTGTGGAATTTAACGTATGCAAACTTTTTCTTTTAGGTGTGGTTAAGTTGCCTCTCTTTT AACTATCTTTGTAAATTGTTTGCTACTTTTCTGACCGGTGGGAGAATTATTAC
>Gm01_5 5645316	GCCGATTAGTCGCCATCACTGGCCGTGTTGAGAAGGTCAGGGACGAATTGGAGCACCTGC TCGACGACGACGATGATATGGCTGAGATGTATCTCTCTGAAAAGCTAGCCGAACAAGATGA CTTGGAGAAGGCGGAGGATACCTCTTCTGCCGACGATGTTGATGATCACATTGACAGG[A/C] JACCTCTTCCCTTCCCTTTTACTCTCCTTCTTTCCCTTCTAACCCTCTATATGTAATTCATCCTT CTATTTTTTTTTTTTTATTTAGGACCGCTCCTGAAATATCGTTGGACAAATGTTGTTGGTAGG GACAGCCATGGAACCTCGCGATAGCGCCACCTACAGCGCCGTACCAAGCAACTT
>Gm02_5 985910	ATTGAAGGAAGTGGAGTATTTGTGGACAGAGACGGTGCAGGAAAGCACATTCAAGCCGGA GCAAGAAGATTTTGATCACAGCTCCTGGCAAGGGAGATATTCCTACCTACGTGGTTGGTG TCAATGCTGGTATTTACGACCCAGATGAACCCATCATCAGCAATGCTTCTAGTACCCT[T/C]C GTCAAGGTCCTTGATCAGAAATTCGGTAATTCATCTTATCCCTACATTTTTTAATCAAACCTT ATTACCCATCTCTACATTATGTGTCAATTATTTATTAAGGAAACTTTGGTCTCAAATTTGCG TGCAAAATTATTATATGGTCCATATTCTCACTGAGCCACCACATGACAGGTAT
>Gm02_5 994292	AACAGTCTCAATTACTTTATATATCATATCATATTAGCATAGCATTAGCACACATATATAGTT ACATACCAGTGTACTTATCAAGATGAAGCTGAATCTCAGAACCACCATTGAGGTATTTGATG TGATCAAAGGCCCAAGTAGGCACATAGTTTCTGCCAAATGGCACATCAATTGGT[T/C]TCTT GGGTTGGCGGCGAAGGAAGCAGAAGCAAGTGATAACAAAATCAGACAAGTCCACAAAGAG GAACCCATTTTTTCTTCTTTGTGTTTCAACTTGTGTTGTGATACGTCCTTCTTGTGCATTCCA ACGGGTATATATAGGCACAAAGACCCTGCTTGGACCTCTTTAATGTAAGGAC
>Gm02_6 754692	AGCCCAACTCACTTGATAGAATCCAACCTGAGTTCCACATTAGACTAACATTCATTCCAACAA TAGAGAAGTTAGCAAAGAGAACAAATTTAGAAGATCTGGCAAGGGTAAATGAGAAGGCTG GACATATCCCAGCATCCTTAGTACAACCTGTCATCAAAGTTGTGGTAGCAAAGTGCA[G/C]GC

	CTGTAAATGTTGTGGCTGCAAGCAGTAATAGACAAATCACATTAAGATAAAAAATTAACA TCAGAGCATGATCGTATAGGGAATTTGCCAATAACAGAAACGTCCATTAAGAAATTGAATAG GGAATTTACCAAACTGAAGCCATTAAGAGGCCATCAAAGCTTTGTTTACAATGA
>Gm02_1 6872184	CTTTTTTGGAGGACTTGAAGGATCACATTGACGAGTTTATTAATGCTTCTATGGATGAACAT AAGACTTGCTTCAAGAAAACAGTTTCAGAAGGTTATTTAGTGATGTACCTAACTTTCTTGTTA GGACAATTTGTGTGCATCCCTTTGTTAACCGATGTAGTGCGGTGTTTGGTTGGT[T/G]TCAG ATGTTTCGGAATGTCCAAGGCTGTTGCAGAGAGGGACACCAATGCTGCCAAAGAAGTTGAG AGTTCTCTGCCTCTTCAGACAACCTTTGCAAGACTAGTATCATGCTTGCTCTCAAGAAAAATA ATGTTTGTCTTCTAGAAGCTCTTTTGGAGGATGTTGATAAAAAATAAGTGAGTGA
>Gm02_1 6872185	TTTTTTGGAGGACTTGAAGGATCACATTGACGAGTTTATTAATGCTTCTATGGATGAACATA AGACTTGCTTCAAGAAAACAGTTTCAGAAGGTTATTTAGTGATGTACCTAACTTTCTTGTTA GGACAATTTGTGTGCATCCCTTTGTTAACCGATGTAGTGCGGTGTTTGGTTGGT[A/C]CAG ATGTTTCGGAATGTCCAAGGCTGTTGCAGAGAGGGACACCAATGCTGCCAAAGAAGTTGAG AGTTCTCTGCCTCTTCAGACAACCTTTGCAAGACTAGTATCATGCTTGCTCTCAAGAAAAATA ATGTTTGTCTTCTAGAAGCTCTTTTGGAGGATGTTGATAAAAAATAAGTGAGTGAT
>Gm02_4 2632633	GTTGGCATGGCCATCCACCAAGGTCACCTTCTATGTCCGCCACGGCCCATGATGAAGGGTT CCGTGCCATGTTGTTATTGGCCTCGCACGTGCCAGGGTAGCCTTTGCTGCTGCTGCTGCTG CTGCTCTGTGTGGCACGTGTCGAGTACTGGGGGAACATGAAGAACTCAGAAAAGGGTG[G/ T]TGAGTCAGAGAAGCCACTACCCTCCTTTGTTCTCTTCTTGCCCTTCATGCACTGCAG AAGCTGCTCCAGCTCCTTCACGAAGTTAATGGTGCCACCAATAATAGAGGCTTGATCACCC TGTTTTTTTTTTTTTTTTTAATAACCGCACATGTGGCACGGTTAGTTGGCTAGCTAGTA
>Gm02_4 2829778	CAGCACCGACGGCGGACTGTGCCTCACGTGAGCGCGTGCCTCCCACTCGCGATCTCGA GCAGCAAAACTCCGAACTGAAGACATCGCTCTTGGAACCTAGATCTCCCGGCGCAAGATA GCACGGGTCGAGGTATCCTAACGTCCCCGCCGGTGGCACGCGAGAATCCGCCACGTGTC[ A/G]CCTCACCGCGAGCCCGAAGTCACCGAGTACGCTTCCACTTTTCGCTGATTAACACA TTGGACGATTTTATGTCCCTGTGAATAACTGGCGGTTTCGGAAGAGTGTAAGGCGCAACCG CTTTCGCGACCTGAACCGCAAACCGGACTCGCGCGGTCCAGCTTGGGGGTCAACCGGTT T
>Gm02_4 2830291	TTCGATTTTCGTTTGAGGTTTTGGGTTTGGTTATTTTACGGCGGCGACGAGAGCGCCGCCG TGGAAGGCGGCCCTTGAGACTGTGCCGTGGCTGCCTTTACCTAGGAAGGTGTCGGCAGAG AAGTTGTTGGTGGCGTCGAGGATGTGCGAGTAGGCGAAGTGTCGCACGGGCTGGGCCT[G/ A]GGCTGGGCTTTTGGGCTTTTCTTCTTCTTGAGGGAGTGAGGGTCGCATGTGGCTATTGC GGACTCGGCGTTGCAAGTGAGGTAGGGCATTGTGAGTAATGATGGAGAAAAGTAGAGAAG AGAGAGCATAACTAATCTGAACACTGAACAGATACATTTTGATTTAAAGGGAATCAAT
>Gm03_2 000300	ATGTGGCTATTCTGTAGTTTATTATGGATACTTGCATGTGGCTGGTTATGTTTGAAGGCCTT GAAACACTTATTTAGAGTTAAGAAGACTGTAATCGTGGAATGTTAGCTGTTAATGTTTAC ATGAACTAAAAAGAGGTCCAGTTTCTCTGGTTGCAAAAATGATAGATTGAAGAT[T/C]GTAAT GCCTATATTTTTATCACGGTCCCTTCTGTTCTTTTGTTCACCTCAAGATAAAAGGAAAAG GATGATGTTTATGGGTGAGAATCATTGGTTTCGTGATTCAATGGTAATAAATCTAATAAGCA AAGTGGTATCGAGTTTTAGTATATTTAGCATAATTACCTAACTAGATGA
>Gm03_9 117708	GCTGGCACGTTTCATCTTCCCTTCTTCTCTTATGTTTTTTCATTTAAATTTTTCGGTCTCAC TCCTCATTACCTTTTCAACAACCTTTTGTTTTTATGGCAGCTCTTAGAAGGTTAGGGGT GCAAGGCGTGGAGATTCGAAAGCCAGAGCAGCTTAACACAATTAGTTCCTCA[T/C]TATCC CTGGTGGAGAAAGCACCACCATGGCTAAGCTCGCCGAGTATCACAACCTGGTCGCTTCTT TACCTCTTTTGTTTTTATTTTATATCTATTCATATTTGCGTAGACTGAAAAATATTTGCCTTT ATCGGAAAGGAATATTTAAAGGGTTATTTTGGTTTGAGTGTGTGGATT
>Gm03_3 5096230	GTTTGTGATCGACTTTATTAGTGATAGTTTTCAATGCTAACTCGAGGCTCCATTACTTTTAT TTTATCATGTTCTGTTAACTAAATTGATTATGTAATTGAGAAATGGATAAAAAACAAATATTGATTA TACAAAATAACAATGGAGGAATGGTGATTATAGAAAAATGCAACACAACT[G/C]ACCTCG ATAATGACATGCATGTCAAATCTTTTGTGTTAAGTCCAATGCCTTCGTAGAAAGCTGTACG TTGGCAATCATTGAGATACATTGTGACATAAACCTGCAGGACAAATTTGACAACCAAGTTTT AAGTAACAGTTAAGAAGGATTAACACAGTAACCTTATGTTGATAGATGTG
>Gm03_3 6398914	AATGATATTTTAAAGCATCTTAGGCTTTAGAATCATGTTTAAAGACTACTGTAGTCAAGTAAAG CTAATAACTCAAATACATACATATTCAGTGACAGAAAACAAGGATTATGTATGTAGTGTC CTGCCAAACAGTTTCCAGTAAAAACAAGCTGCATGTTTCTTATTTCAAAGCT[T/G]ACAGG AAGGTGATAATCGTGCAGTACACAGCATACAACCTACATAAAAGTTTCTTATGGCATAGAAA ACATGTAGTCCTCCTATTCTGACACAGTCAAGTTTAACTAAACAAACAAAGAACTCAAAGA

	<p>           TGAGAGTATAGGAAATACAAATACAATACTAGAGTCATTCTGTACAAAAGG            AAGTAAAGCTAATAACTCAAAATACATACATATTCAGTGACAGAAAAACAAGGATTATGTAT            GTAGTGTCTCGCCAAACAGTTTCCAGTAAAAACAAGCTGCATGTTTCTTATTTCAAAAGCT            CACAGGAAGGTGATAATCGTGCAGTACACAGCATACAACCTTACATAAAAGTTTCT[A/C]ATG            GCATAGAAAACATGTAGTCCTCCTATTTCGTACACAGTCAAGTTTAACTAAACAAACAAAGAA            ACTCAAAGATGAGAGTATAGGAAATACAAATACAATACTAGAGTCATTCTGTACAAAAGGTT            TGGGTTTCATCATGTAATAAGACAATACACATACTAAGACACATAATTCTCTTCC         </p>
>Gm03_3 6398969	<p>           CCTGCCAAACAGTTTCCAGTAAAAACAAGCTGCATGTTTCTTATTTCAAAAGCTCACAGGA            AGGTGATAATCGTGCAGTACACAGCATACAACCTTACATAAAAGTTTCTTATGGCATAGAAAA            CATGTAGTCCTCCTATTTCGTACACAGTCAAGTTTAACTAAACAAACAAAGAACT[G/T]AAAG            ATGAGAGTATAGGAAATACAAATACAATACTAGAGTCATTCTGTACAAAAGGTTTGGGTTCA            TCATGTAATAAGACAATACACATACTAAGACACATAATTCTTCCCGCCCATCTCTCCTCGT            ACTTTCCAAATGAGTAGGTAGTTGAAGGCATCCCAGTCAAGTGAAAGGGGTC         </p>
>Gm03_3 6399038	<p>           GATGGAAGGTTCTTGCTTTTGTGTATATATACATACTATTATTGCTAGGAAATTGAAGACC            TAAGATACAATAGAGATGGCAAAGCTATTTTTGAAACAGGCAAAGCAATACGCAGATGCAA            GACCAAGCTATCCTCCACAACCTTCCAATTCTGCTTCCAAGACTCCCTCTCA[A/G]AACCT            TCGCTTGGGACGTCGGCACTGGGAGCGGCCAAGCTGCCAAATCTGTAAATTTTCTTTCTTC            TCCTTCTCTTTCTTAATTCTTTACCTATATATATATATATATATATATATTTCTTGGTCAATTTCC            ACAAGCACTCTCACATATTTTACACGGGCGTGAGATCCTGCTTAACCTAG         </p>
>Gm03_3 6416992	<p>           AAATGGGTTGATGACAATTATAGAAGCATTGATTTTCCATTTGAGCCCGTGGATGGAGCTGA            TCACACAGGACCCTTTGAGTTTGTGACGGAAACAATGATGGATTTGGATGATTTCTTGACCT            ACATAAGATCATGGTCAGCATATCAGACGGCTAAGGAGAAAGGAGTGGAGCTTCT[G/A]GC            GGAGGATGTGGTTGAAAAATTCAAGCTTGCTTGGGGTGAAAGATGCTAAAAAAGTTGTCAAG            TTTCCAATTTATTTGAGAATTGGAAGAACAGGGGATTCTAAAGACATATGCAAATGGTTGC            TTTTACTGTGTGGGAGATGTGACGAGTACCAACTTTTATGAGTTTATCCATTGAT         </p>
>Gm03_3 6418097	<p>           GACATTGCAAATTAATTTTCCATACAAGTTTCTCAATCCTAATATAGTGATAGTGATAAA            TAGAATAAGACAACAAATTAGTACTAGGCTCTGTCAATCTTCATAGGCCAATCTTTTACC            ACTCTTTAATAACTTATGACCCGATATTTGAAGGTGAGGATAACAAATTGTAACA[G/A]GAAAA            TTACAGGAAATACGGATTTTATTACATGCACGTAAGAAACACATCAAATTTTCACTCTCAAG            AATCCAAAGAAAAGATAACAATGCATGCTCTCAAATCCTCAACATGCTAACTAACATCTTGC            CAATGCAACTTCTAAGGATGAATATTTAGACAAAAAAAACATTAAAGAG         </p>
>Gm03_3 6460374	<p>           CCCTTACATATTTTACCATAATAGTTTAGTATGAAGGTCCTTTTATTTTATGCTGAATTAATC            CACACCACTAACCACAACCGCGGTTCCCACTTCCAGAATCCCCTCTTCTGTTCTTGATG            GGTTTCCTTCATCACCTCTTCCCTGAAGCCACATAACCCCATGGCAATGGCAC[T/C]JAGAC            GCAGCACTCTGGATCGTGAAGATGACATGGATTGCTCTCAGTGGCTGGATTTCTTCTTGTT            AACTGTGCTGACGAATTCGCCTCTTCTCTACGCTCTGGCGATATTGGCCCCTTTCATGTG            GGTTGATCCATTTACGCTCTCTATTTCTCCAAGATGGGTACTTTACTTACTG         </p>
>Gm03_3 6468024	<p>           GTATGAAGGTCCTTTTATTTTATGCTGAATTAATCCACACCACTAACCACAACCGCGGTT            CCCACTTCCAGAATCCCCTCTTCTGTTCTTGAAGTGGGTTTCTTCTACACCTCTTCCCTGA            AGCCACATAACCCCATGGCAATGGCACTAGACGCAGCACTCTGGATCGTGAAGAT[C/T]JAC            ATGGATTGCTCTCAGTGGCTGGATTTCTTCTTGTAACTGTGCTGACGAATTCGCCTCTT            CTCTACGCTCTGGCGATATTGGCCCCTTTCATGTGGGTTGATCCATTTACGCTCTCTATTTT            CTCCAAGATGGGTACTTTACTTACTGGGTATGCAGCAAGTTTTGTAATTTTCTGA         </p>
>Gm03_3 6468052	<p>           TCTTCACGACTCATATGATACTCGTGACAAAAGACATACTCAGCAACAACCTTGAGGAACATT            AACAACCTGGCTCCTTAATTTTGGTGCACCAGCAAGCACTGCAACAACCTTTCTGATCTTTGC            CATGCTCTGTGAGCAAAGGCAAGTGGTCGTGCTCACCCATGTCTAGATGCACTATG[G/A]GC            ATCAGTGACATCAATCGCTCTTTGATGTGAACTCTCGTCCACTCTACAACCACCAATTATCA            TAGGTGAAGTCATTTTGTACATGAAAGTTAAAAAATAAGATGTTAGGCCGATTTTATTTAATG            TAATTGTGTGTAATAAAAATACATATAAATGCAAATACATTTAATATACTTTTT         </p>
>Gm03_3 6553944	<p>           TGTTCAGGAGGTTGAAAAAACTCATAGATTAGAGTCCGAAATTCTAAGTGAAGGCTTTTG            TCAATATATGAGAAGCAAAGGATTTTCTAATCAAACGATCTTAAGTGCTGTGATGTTAAATTT            GACACCTAAAAGAGTCTTCACATATTTGTCATGCATGAATGGCAAGTTCGAAAA[G/A]TAGA            ATGGAGTAATCACATCACATGCCGCTTCTATATACATATGATTTGGTAAAAAAGCAAGTGGC            CGGCAGGAAGGTGCATGGATTTTAGGAGGAGTTAAAGAAAAGAATATTGTATAAACATAAA            TGTTACAATCAGCCTCACGTAGCAAAGTGTGACACCCTCTTTTCTCACAATA         </p>
>Gm03_3 6559857	<p>           ATGAGAAGCAAAGGATTTTCTAATCAAACGATCTTAAGTGCTCTGATGTTAAATTTGACACC            TAAAGAGTCTTCACATATTTGTCATGCATGAATGGCAAGTTCGAAATTAGAATGGAGTAA         </p>
>Gm03_3 6559926	



	TCACATCACATGCCGCTTCTATATACATATGATTTGGTAAAAAAGCAAGTGGCCG[A/T]CAG GAAGGTGCATGGATTTTAGGAGGAGTTAAAGAAAAGAATATTGTATAAACATAAATGTTACA ATCAGCCTCACGTAGCAAAGTGTGACACCCTCTTTTCTCACAAATATATGTACTAATAATGA AAGGAATAGAAAGTTAGAATTAATGAAAAGTTTTAAAACACATTTAAATAAAA
>Gm03_3 6560002	CATATTTGTCATGCATGAATGGCAAGTTCGGAAATTAGAATGGAGTAATCACATCACATGCC GCTTCTATATACATATGATTTGGTAAAAAAGCAAGTGGCCGGCAGGAAGGTGCATGGATTT TAGGAGGAGTTAAAGAAAAGAATATTGTATAAACATAAATGTTACAATCAGCCTCA[A/G]GTA GCAAAGTGTGACACCCTCTTTTCTCACAAATATATGTACTAATAATGAAAGGAATAGAAAGT TAGAATTAATGAAAAGTTTTAAAACACATTTAAATAAAAAGCATTTCAAAAAGGATAAAAGGTT TACATTCGTTTTTCTAGCATTCTAATAAAACTTATTTAAATAAATAATAAAA
>Gm03_3 6612938	GGAGGATCCGGATCCGAAAGAAGGCGGGTCCGGAGCGTGAGACTAAAGATGGTTCCTTTTT ACGAAAGAAGCGTTTGGGGCTTAGTTTGAGGTTTTTGGGGTTGATGGTGCTAATGAGCTTC ATTGTTTTTGGAAAAGAAGAGAAGAGGAGAGTGGGGTTGAGGGGAAAGAGAGTGAAA[C/ A]GTGGAGGGATTAAATGGGGGCACCATGGAAGGTCCATTTCTCAGCCTTGCTACCTTCCA CGAAGTTTCTATACATCATGGACTCATTATAAAAAGAAGGTGGCTATTTTTATTTTTATTTTT TTCTTTATTAATTAATATGTCATAATAGTGATTGTATTGGTTTGAATGCCTCAT
>Gm03_3 6634361	CAATATGGAGGGCAATAGGAACAAAATATTAATAATGTCACTGGAAAAGAAGTACTGATCTTGAC ACTAAGTGTCAACAATGCTTTGATCCATACCCGAATCTGTGCCCGAATGATGATCAATGGAA TTGTAATCTGGCATGTCAGTTAAGGACCTGGCTGGCATGGATTCTTCATCATCTC[T/C]GTGT TCAGATGCAGAGCTAGCAGCATCTATCTCCTTCCCTGGACACTCATCTTGACACATTTCAG CAACTTCAATGTTGTAATTTTCAAACCATGACTGCTACTACTTAAGGGTGACAAGTCTTCTT TCTTGACAACATTAAACACAATCCCATTATGGATCCTTTCAGGCTGCAGAAG
>Gm03_3 6808892	TGAGTCCCAAGGCTTCACTTCATTTCCACAGTAATGTACCTGAATCTGACATTCCTAAAAGT TTCTGACATTGTCTTGACCCCATCTGTTTGCGTTGTCTTGGCTGGAAGCTACATATTGGAGG TCGATGTTAGGAAGATCTAGCAGCAACTCAATGTTGGAATCTCTCAGAGCTTGGA[G/A]GAC TTCATGGTTTCGATCATAGATTCTCATCCTTTGGAAGCCGTAAGTGTGTTAAAAAGGGAAACTA CTTCTTGTTGGCGATGGCAGATTGTTGCCAACTCTTCCATAGCATAACCCAGATTGTGCACCT GCACCAGATGGAACAAAATGGTGAATTGATAATACTACTTCTAAGATATCAAG
>Gm03_3 6952394	AAAAGAAATAAAAAAAAAATAAGATCTGTTTTTTTTTTTTTTGAGAAAGATCATTTTTTATTATT ATTGTTTTGAATTTTGAGAATTATGTAATGATGAGAGGTTGATGTTGTGAAAAAATAATGG AAATCCAGGGAGGAGGCGGTGGTGGAGGTGGCAAGAGTGGAAAGACAATGT[T/G]ACCA ACAGCAAGGAGAATTTGTTTCATAGTTTTTTCTTGTTACACTTTGGTACTCTTCAAACATTGGTG TGATCCTTCTGAACAAGTATCTGCTCTCAAACATATGGATTCAAGTTCCTCAATCTTCTCACA ATGTGCCACATGTGCGCCTGTGCTGTCTCAGCTATGTCTCCATTGTGTT
>Gm03_3 6960275	GTTAGGCAATTCATGTCAATGTTTCATCTACTTGTTTTTCGAAAGGTTAGTGTCTTAAAAACCA AAATACGAAACCAAAAAATGTTTCATGACCCAAAAACAAAAGAAAGGAAACGAAACCATAC TCTTCTAATATTTTATGCCAGCTCCGGAACAGATGGCAGTGCCTCATAACCATC[C/T]CGGT GCTGCTTACTAGCATCTCTTTCAACTGGGATGCACTTGAGGATAGCAGCTATAGGCATGCT AAATGCTCCGATTACCACACTTAGTAACCAGAATTGCCAGTTTAGAGGCACAGTGCTGGCA AACGTTCCAAGGAATTCAACTATCACCACCTTGAAATGCAGCGGTGGCGAAAATT
>Gm03_3 6996777	CATATATTTGCATTCACTTAACGTAGTCTTGATACCCCAATCCACTAACAATATTTTTTTTATA TTATTATTATTATCATCATCATCAAAATGCCTGGCTTGTAATGAAATGGATTTGGACCAAAC TATCTAGTGACGTGCGCAATGAAATCATCCAGTTCGCGCGAGAAAACACCCC[A/G]TTCATC CTTGGACCTGCGAATAGCATTTTTCAGGTTTCATTGCCCTTTGTCTCATCTCATCACCTTCTTT TGTTGCCATCAATCTCCTCACGGCATTCTCAACATCTGATGATGTCACCAACTCGTCCCTGT GGTCCCAGTCCTTCACAACCACCCCAACTCTGAGCACTTCTGTTACCAA
>Gm03_3 6997184	GCACCCCATTTGTCATGCTCTCCATGCAAGAGTTCCATCCACAGTGACTCATAAACCACAC AGTTGAAGTGTGACTTAGAATCTCCAATTGGGGTGCCAGTCTCTCACAACCAACCCCGTG CCTTTCACTCTCTCTTCAAACCCCTTTGGAAGCTCAGCAGTTCTCACGCCATCCTCA[C/T]G GAACACATCTCCCTTGTGAGCATCCCTCACCACCCATATGAACTTTTGCTTGCTTTTTTCCA ACCCGTTTGCAACCTCCTTGATTGCTCCTCAGAGAAGCATGTTGTTGTCCCAAACGACACA TACAAAACCTGACCCTGCTTCTTGCTTGTAAGCCACTCAACACTGAAGTGCTTG
>Gm03_3 6997185	CACCCCATTTGTCATGCTCTCCATGCAAGAGTTCCATCCACAGTGACTCATAAACCACCA GTTGAAGTGTGACTTAGAATCTCCAATTGGGGTGCCAGTCTCTCACAACCAACCCCGTGC CTTTCACTCTCTCTTCAAACCCCTTTGGAAGCTCAGCAGTTCTCACGCCATCCTCAT[G/A]GA ACACATCTCCCTTGTGAGCATCCCTCACCACCCATATGAACTTTTGCTTGCTTTTTTCCAAC CCGTTTGCAACCTCCTTGATTGCTCCTCAGAGAAGCATGTTGTTGTCCCAAACGACACATA

	CAAAACTGACCCTGCTTCTTGCTTGCAAGCCACTCAACACTGAAGTGCCTTGT
>Gm03_3 7024877	TTTCTCATTGTTTTAAAAGGGTGTTCACTTGTCCTTTTCCATGAATTGCAGTTTATGAGCC ACAAGTGCAAATGGGAATGAAATGAAGAATGGGGTTTAGAGTTAGTCAAGACAAGTGGTGT GGTGTCTATGTTATGACTGCACACATGTGAAGTGAAGTAGAGACTATTCAGTCC[G/C]CAG CAGCTGTTTCTAGTGTGTGTGTCATTGCATTCTCATCCTTTTCTCTTTTTTACGCTTAA TTTCTTTCTCTCTTTCTCCCTCTTCTCTCTGGAATTTGGAGCATCAGCCAGCACTCTATGG ATTCTCTGATTGGTAATTGGCCATCCTACGATCCTCACAACTTCAGTCAGCT
>Gm03_3 7024958	GAAATGAAGAATGGGGTTTAGAGTTAGTCAAGACAAGTGGTGTGGTGTCTATGTTATGAC TGCACACATGTGAAGTGAAGTAGAGACTATTCAGTCCACAGCAGCTGTTTCTAGTGTGTGT GTCATTGCATTCTCATCCTTTTCTCTTTTTTACGCTTAAATTTCTTTCTCTCTT[G/T]CTC CCTCTTCTCTCTGGAATTTGGAGCATCAGCCAGCACTCTATGGATTCTCTGATTGGTAATT GGCCATCCTACGATCCTCACAACTTCAGTCAGCTTCGACCTTCCGATCCTTCTAGTTCTTCT GTAAGTTGCTGTTGTTGTTGTCTATGAAATTGATAATCCTGGTAATAATTACT
>Gm03_3 7027198	GTAAAGCAACCTTTTTGTAGAAGGGTCTTCTATAATAACTCTAGTTTCATTTTCTCCAATGTT ATTTTTCAAGTTGCTTAACTTTTGTTTTTTAATTTTCGGTAGACTCAATGGTGTCTTAATTC AAAGTTTTGCTAACCTGTTGAAGTTCCAACAGTGATAAGTACTGAAGCCAAA[G/A]CATCCTC ATGAGACATATTTATCAGCATTCTGAGCAGAAGGTTAGTGACTTGATGGTTAAAGAGCATGT GTTTTGGTGCAGTTAGGTATGCATATGCTTGATGCTCATAACCTTTTGTGTTTTTCAGTTGAA TCCAAAAGAGCTGCATCTGATAACCTTCTTTACCAGAGCATGGATG
>Gm03_3 7144398	TCCTTAAAAAGTTGACACCAAACTGAAAAAGCATTATCTATGTTGCAATGTTTAGGGCTT TCGTAATTTGCAACAAGCAGATGGGAGTTTTTCAATGACAGGTTCAAGAAAGGTGAAAGA GAACCTTCTACATGAGATTTCGGCGAAGAAAAGCTTGGACCAGCAAGCAACAACCAA[A/T]GG CACCAAACCAAGCAACACTCCAAGACTCTGATGAAGATCAAAGGTCCTCATCAATCTCATCA TCTTCTGGTTACACTACCCTTGTGGATGAAAAACAAGCGACTCAAGAAGGAGAATGGGGTGT TGAACCTCTGAGCTCACAAGTATGAAAAGGAAGTGAAGGAACCTCTTGATTTGGT
>Gm03_3 7165409	AAGATTCACACGTACCATCTCGAAACAGTTGTTCTGATAACTGTTCTGAGAAGCTCCATCA CAGACAGGTGTACCAGAGATTAGTGTTTTACTCTTTCTTTCATTGTCAAGCCAGCGCCCAAT TAAGCCTATCATTCTCGTGTCAAGCTAATCTCCGCAATGGATGTATTTCCTGAT[G/A]TGAC GGAGTCAGTGCCCCCAATGCAAATTTCAAGGTCATTAAGTAAGCTTCTAGGATGAGAAGGG ACATCAAGAAGTCTTTCTTGAATAGCTTGGCAGAGGTTTTGCAAGTCTGCAGAACTCACTTC AGTAGGTTTGACCTGGAAACAGGGAAACAGAGAATTACATATTTGAAGATGCT
>Gm03_3 7300009	GCTCTCATCGATGCCCAAACAACCCTAGTTGCGAATTTAGAAAGGAACTCCTCGCTGGCG CTCACCATTTCCCTCGCGTGCTGCCACGTGGCGTCAAGGCGATGAGGACGAGGCCAGAG GAGGGGCGGAGGGAGGAGATGTGGAGGGCGGAGGCGGAGGAGGGGAAGAGGTAGAGG G[C/T]GGGGGGAGAGCGGTGAGGAGGGGGGAGAGGCCGCGGCGGAGGCGGCGGGCGG TGAGGGAGGAGGCGCGGAGGAGGGGACTTGGTGAGGATGGGGGTGGTGGAGAGCTTGTG GCGGGCCTCGTGGGGGTGTTGGATGATGAGGATTTGGGTGGTGGTTTGGATTGGAGGAAG TGGAAAGCGC
>Gm03_3 7730768	GCAGGTGTGTGCACCTTGTGTGCTGAGCTTCCATGCTCTGGTGCAATGGAATAAAGTGACA CTTCAAGAGCAGCAACTTCTCTAAGCTCCTCTTCGAGCTTTTCAATTCGTAATTCATTTCTT CAATTTTTAATTCTGCAGCTGTATCATCTCATGTTTACCATTTTCTTACGATT[T/C]ACTGT TTCATGAATTTCTCTTCACTTCTTACACTCTGAGCACCATTGGAGGATCCATCCACAACAT CAGGTTTCTTGACCTCTTGAACAGGTTTTTCTATCTACACCTTCAGAAGATTCTGATGATACA TTTACAGGAACTTTACAGTCTTGAATTGGTTGAGGGAGTATTTTTGTTT
>Gm03_3 7731694	CCTTGAAGCTTTTCTCTTCTCAATTTCTTTTCATGCTTTCAAATTCCTTCAAATATCAATCAAAT AAAGATGATGTGATAGGGGGAAATGATAGCAGTAACTAACCTTAGCGAGCTCCTTAAGCCT TAGTTCTCAAATCCTAAAGTCTGTTTGTGTAATCACTAACAATACCTACTCC[G/T]TAGAAT AGTGTGTTTGGCTTCCACCACGAGAGAAGTGCTTGTAACCTCTCTATCTTTCCAGAGGCTC AAAACAACAATTGTCTACCTGCAAAAGTATTGGTAAAATAAGTGGCTTGATTCTCAGAGAAT AATAGACAAAATCGTATAAGGAACCAATTTTGATTATCGGGATAAGAG
>Gm03_3 7787006	TATCCCTGAAGTTTATGCTAAAAAGCAAGTCATTGTTGCATTCCATATTGCTCTAAACTGCAC TGAACCTTGATCCAGAGTTGCGACCTAGAATGAAAACCGTGTCAGAGAACCTTGATCACATC AAAATACAGTGAGAAACAAGAGATGTCTTAGTGTAGTGGGGTTGATTTTGTAACT[T/G]GCG GGCTAGTAGATGTATCTTTCTTCCATTTCTCAATGGAATTTGTTTCTTTTACCTTTATTTA AAATCAAGGAACGGCCCAAATTGATTGGTGTGGTAAATATTTTCTTCTTTTACATCCTGTT TACTTCAAATACAAATCCTACGGATTGCGATTGAACACCTTGCTTAAAT
>Gm03_3	GTCTAATTAGTATTATTTCTTTCTTAGTTTCTTCTATATATGCTATCACATAATCAACAAA

7823922	AAAAAAACACAAACACTTGATAGTACTACTCAAGGTATGGCTTTTCAAGTGTTGTTCAATTTGT TTGATCTCAACCATCGTGTGTTGCATCCATACTATGGAGAAAACAAAACAAA[C/A]CAAGACC CTTCTTCCACCGAGTCCAATGCCCTACCCATCATTGGACACCTCCACCTTCTTTCTCCAAC ACCTCACCAAGATTTTCACAAGCTCTCACTCCGCTATGGACCCATAATACACCTTTTTCTTG GTTCAAGTCCCTGTGTGGTGGCTTCCACAGCAGAAGCCGCCAAAGAGTT
>Gm03_3 7825579	GAGTTTAGGCCAGAGAGGTTTGTGAAAATGGGAAGAGTCAATTGGATGTTAGGGGACAAC ATTATCATCTACTTCCGTTCCGGTAGTGGAAGAAGAGCATGTCCTGGTACTTCTTTGGCATTG CAAGTTGTGCATGTGAATTTGGCAGTTCTAATTCAGTGTTCCAATGGAAGGTTGA[T/A]TGT GACAATGGCAAGGTGAACATGGAAGAAAAGGCTGGCATCACTCTTCCGAGGGCTCACCCC ATAATTTGTGTCCCTATTCGAAGACTTAACCCATTCCCTGTTGTGTGATTGTTTTTATTTTA AACATCATTTTTCCATTACGTATATGATATGATGTCATGTCAAACATTTATGC
>Gm03_3 7827399	GTCGAGTGATCCATACCCGCTAGGTTTTACACATTCTTTCCCTGAAACAAACACCACCCCT TTTAGGTCTTCTTCTAATCCTCCCTCTCTCTAATTCACACGCAGCGGCTTCTTCTCCGTT TTCCCTGAATCGGTGTGAACAAATGGCGACGTACGCTGCGATGAAACCCACCAAGC[C/G]AGG GTTGGAGGAGTCCCAGGAACAGATCCATAAGATAAGGATCACCTTTCTTCCAAGCACGTC AAAAACCTCGAGAAGGGTATGCACCTATTTATTCTACTTTTTGTTTTTAAGGTTGTTGCTATC CTTAGCCTTATGTTTCTAGTTTTGTGTTACTGTTAGTTTGTGCGGACTTGGTT
>Gm03_3 7828684	CAACACATGGGATAGATTTGAACTTCGTGTCCACAAGAGAGTGATTGACCTCTACAGTTCC CCAGATGTGGTTAAGCAGATTACCTCTATCACAATCGAACCTGGTGTGGAAGTTGAGGTGA CCATTGCAGATGCTTGATCTATATCCAAATTTTAACTGTTATGGTTTCCTAGTTTTT[G/A]GTC TCCTCTTATTCTGTATTTTATAAGTTTTGATTGCGGGACCGAGTTGAATTGTTTTCTTTGTTT TACGGTTGTAGACGGTTGATGAATACCTGAGAAATGTCATTTTTATATATTCTAGAATCCTAC ATATATTTGTTTTGAAATTATTACGAATCATTTCTAATGGACTCAGCTAA
>Gm03_3 7828746	CAGATGTGGTTAAGCAGATTACCTCTATCACAATCGAACCTGGTGTGGAAGTTGAGGTGAC CATTGCAGATGCTTGATCTATATCCAAATTTTAACTGTTATGGTTTCCTAGTTTTTCTGCTCC TCTTATTCTGTATTTTATAAGTTTTGATTGCGGGACCGAGTTGAATTGTTTTCTT[G/T]TGTTT TACGGTTGTAGACGGTTGATGAATACCTGAGAAATGTCATTTTTATATATTCTAGAATCCTAC ATATATTTGTTTTGAAATTATTACGAATCATTTCTAATGGACTCAGCTAAGATTATATCTTTTC CTTTTCTTTATCATGTTCTGCAAATTTTAAACCTGAAACTGATTGT
>Gm03_3 7832228	CCTTGTGTTTCATGAAACTGCCTTTTGTGTTTTAGGTACCAACACATGGGACAGATTTGAACTT CGTGTGCACAAGAGGGTGATTGACCTCTACAGTTCCCCAGATGTGGTTAAGCAGATTACCT CTATCACGATTGAACCTGGTGTGGAGGTTGAGGTGACCATTGCAGATGCTTGATCT[C/T]GA TCCAAATTTTAACTGTTATGCTTCCTAGTTTTTTTTCCCTCTCTTTTGTATTTTGAAGTTT GATTGTGGGACTGAGTTGGATTGTTTTCTAGTTTTACGGTTGTAGACATTGATGAATACC CGAGAAATGTCATCTTAATATATTCTAGAGTGTTACATATTTTGTGTTGAAAT
>Gm03_3 7863675	CTTTGACTAATGTTTGCAAGAGCATCAGTCCCGCTAACGGTTGCAGCATCCGAGTGCCAC GCTTCGGGAAACGCCTCATCAGCTTGTTGTAACCATCCAAACAGCCAGCATCAGCTCCCCCT TCGCCTTGTCACCCCTTTCTGTCTCCAGCCTATACCACTGCGGTGCCAAGGGACTGT[G/A]T GGTGGAAACACGCTTTGGGATCTCATTGAGGTCAAATAAGACACGACCAATGAAGTCATCTT TCACGACATCCTTGTCCTTCACAGTGACCTCCAGTATGGAAGCCTGAATGCGGTCTTTGGA GAAAGCAAAAACCTGGTTCCATTGCGGATTAGACTTCTGTGCAAGTGCCGAGTGGT
>Gm03_3 8055240	GATATATATTGACTTCCATGTTTATTTTCTTATTAGTTTTAGTGTAACATAGTATTTATACA TTAGTTAAAAATTGGTTTGACAAGGATGGAAGTCAGTATTATTGGTTTGGTCTCAACCTGG CTGGTTCAAGAGGAATCAATCATCTGATGGTAGAATCTCACTCTTTTAGTCAT[G/A]AAGATG ATAAAGGATGGCTGCTTAGACACCCACCTGAGTCCGCCGACTAGTCAAATCTGTTCTTTC TTGGAATCACGTATTATTAGAGAGGCCAACCAAGTAGCTGATGCTTTATCCAAACATGGTC TTTCTTGGTACTACTGATAGCATTGTTTTTTTTTTTATTTCAGTTCCGTCT
>Gm03_3 8055290	ATAGTATTTATACATTAGTTAAAAAATGTTTTGACAAGGATGGAAGTCAGTATTATTGGTTT GGTCTCAACCTGGCTGGTTCAAGAGGAATCAATCATCTGATGGTAGAATCTCACTCTTTTAG TCATTAAGATGATAAAGGATGGCTGCTTAGACACCCACCTGAGTCCGCCGACT[G/C]GTCA AATCTGTTCTTTCTTGAATCACGTATTATTAGAGAGGCCAACCAAGTAGCTGATGCTTTA TCCAAACATGGTCTTTCTTGGTACTACTGATAGCATTGTTTTTTTTTTTATTTCAGTTCCGTC TTTTCTTTCTGTTAAGATATGTGCTGATGTATATTGTCTACGACCTTCCCT
>Gm03_3 8065066	TGTGGAACCACCACTCAAGACTATGTTACCATAGAGATCCTTCCTGATGTGACATCACACT TCATGATAGAGTTATATGTTGTCTCATGGATACCAGGAGATTCATCCCAATCATGGATGGC TGGAAGAGAACTTCAGGGCACCGGAATCGTTCAGCACCGATCGTAATCACCTGCC[A/G]AT CAGGTAGCTCATAGCTCTTCTCAACTGCTGAGCTGGTCTTGGCAGTTTCCAACCTCTGCTC

	ATAATCAAGAGCAATGTAGGCCAGTTTCTCCTTCATGTCCCTCACAATTTCCCGCTCCGCAG ATGTGGTGAAAGAGTAACCACGCTCAGTCAAGATTTTCATCAGGGCATCAGTGAG
>Gm03_3 8083417	GTTTTCTTTTCTCTTCCATCCATGGATTCTCTGAGTTCCATCAACGGCGATTTTCGGCTTCGA CGGCGGCACGGATCGCAGATTCGCGTTTTTCGCGCCAAGCCTCCTTCCAACAACCTCACAC GCCGATAGACATTCCGGCGCACGGCCACTACGACCACCACCTACTGGTCCCCGCG[T/A] GACGATAAACCCCTCCGCCGCTCTCTCCAAATATCCTCTCTCTCCTCCTTCGTTTTCTCCGT TTTCCGAAGCGTTAGGTCCGGCCACAGGTACATGAAGAGGCTCTTTCTCATGATCTCGCTC AATGTCGCGTACTCCACCGCCGAATTGCTCACTGGACTCTTCACCGGTGCGGTAGGT
>Gm03_3 8117453	GTACATGAAATTATAATAACCGTTTTCTGCACCTGTACTTAATTATTGGCCTTACTTGCTATT CTCATGAGCTACAGCAGCAATGTGGTTGCTTGTGGCCCCAGCAAAGTCTTATGGAACGC GCTTTCACCATGCCAAGCTTCATCCTCTTCTTCGTCGACGTCATTATTATCCCTG[G/A]CTTC TTCTCCTCCTTCTCCCCAAAGGGCTCAGCATCATCAACACACTCAATTTGCTTCTCGAGAGA ACAAAAAGAAGGGGAAGAAGAAGAAGGGTGAGGCACGTGGTGCACGTGCACCTCCTTTGG TGCAGTTGATGCAATAATGGAACGGCTTCTAACACTTGTCACTTGTCTGAGTCT
>Gm03_3 8117485	TTGTACTTAATTATTGGCCTTACTTGCTATTCTCATGAGCTACAGCAGCAATGTGGTTGCTT GTTTGCCCCAGCAAAGTCTTATGGAACGCGCTTTCACCATGCCAAGCTTCATCCTCTTCTT CGTCGACGTCATTATTATCCCTGGCTTCTTCTCCTCCTTCTCCCCAAAGGGCTCA[A/G]CAT CATCAACACACTCAATTTGCTTCTCGAGAGAACAAAAAGAAGGGGAAGAAGAAGGGTG AGGCACGTGGTGCACGTGCACCTCCTTTGGTGCAGTTGATGCAATAATGGAACGGCTTCTA ACACTTGTCTGCTTCTGAGTCTTCCCTTCCCCAAGTGATGCTCACACAGCGAGT
>Gm03_3 8118175	TCCCATCTACAACGTATGCAACACCAACACAGTAAACAAAAAAAAAAAAAAAAAACACCAC CAGCAAGCAAGACACAGTTAGTTAGTTAGTTGCATTACATGCATTGAGTTTTTGGCATTAC CTTGAGTAGAAGATGAGCTTAACGGTGGCAACAACCTCAGCTTGTGTTTTTCCGA[G/A]TCA AAGGTGCGTCCTTCTCTACAAAACAAACCAACCCAGAAAGACCCACAATTCCATAACATAC ATTTGCACATACTTGTACGTGAATAATAAAATTTACGGCAAAAGGGTGTGAATTAAGTGA AAGTTGGAGTGTAATATTTTGAAGGAACAAAAACGCAGTGATGTTGGGGACCAG
>Gm03_3 8119584	AGGTTTCAGATTTCAAAAACAAAAATAAGATAAAGAAGAAAAAGGGTATCAACAAATAAACCA AAAATCTATAGCTACTACATCTACATGACCCTCCAAAGACCAAGAAGAAAAATAATTATGTT TTTATCAACAACCTTTTCTCTCTCTCTATCTCTCTCTCTCACACAAACGCAT[C/G]AATACA TACACTCTTTTCTTCTCTCTCTCTCTCTATCTCTATAGCACTCCATATATATTGCTTCCATGA GTTATTATTCTCTCTCTAACTTTTATTTGTTGCACCACAGAAAGGGAGTTGTATGGGTGAGG ACGGCACGTGCGATGGAGTTAAGGGAGGAGGAGGTACGTGTGGAACCTA
>Gm03_3 8131888	GAGGCATTTACTTTAGAGGGCATAACATAACTGCATATTATATATTAATTATTTGCAGGGAG GAGGTCCCTTCATGGAACGTACTACTAGGAAGAAGGGACGGTTTGATTGCAAACCAATCTGG TGCGAACACATCCATTCCCAATCCAACAGAAAGCCTGGCCAATGTCACTGCCAAT[C/G]TG CTGCCGTTGGCCTAAACATAACTGATCTCGTTGCATTATCTGGTAATCATATTATATTGCTT CCCCGTCTACTTTCTTGTGTGCACACACAAACTTCTATCAACATATGTAGTGAATAAAAA ATAATATAAAAGAATTTTAAACAATTCATGTCCTGAAATCAATTGAATTAGAT
>Gm03_3 8132996	TGACAATTTACCTATAGAAAAATGGTTGATTTAATTACATAATTACTATGATGCTTATTAAGTA GTGGTTTTGTTTGAGTGGAAGGTGCACATAGTTTTGGTCGTGCGCAGTGATGATTTTTTC AACCAGAGACTGTTCAACTTCAGCGGCACAGGTAGCCCTGATCCCACGCTGAA[C/G]ACAA CCTATTTAGCGACACTTCAGCAAAACTGTCCACAAAACGGAAGCGGCAACACTCTGAACAA CCTCGACCTTCGTCTCCGGACACCTTCGACAACAACTACTTCCAAAACCTTCTCAGCAATC AGGGCCTTCTCCAAACAGACCAAGAGCTCTTTTCCACCAACGGCGCGCCGACACA
>Gm03_3 8136442	TCCATCACATCACATCACAACACAACATAACATTCATATGAACCAGGTGAAGCTCAAGTTCT TCTTCTTGTTCCTTCTTCTCATTTCCTTCCATTTCCATTTTCTCTATAAACCTCTCCTCCCTCT CTCTCTGACTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTAGATC[A/G]CACCGC ACCGCACCGATGGAACACAAATGATCCAACCTAACCCAAGCGCCTCTTACCTCCAAGGTC TTCCATTTTCTTCTTAATTCCTCTTCCATATCCCCTCTTTTCTTCTCTCATCTATTTATTTTCT TTCTTTTAATTGCATATATGTCAAAAACAATTTTCTGCTTTTTTTTTT
>Gm03_3 8150181	ATATATATACAGTGGGTATGGAACGTGTGACATCAGAGATTCCGGTGAGGGTGGTGGATGA CACCGACGATGGTGCTGCACTTTCATCTCCAACAGTGCAAGTGCATCTTTTCAGATGGATT ATTGTGTCAGGAACAATAGAAAGTCTAGTGAAGGTGCCAGCGATGACGACGAAAAT[G/A]G CTCATCAAGGAAGAACTAAGACTCTCCAAACAACAATCAGCTTTCCTCGAAGACAGCTTCA AAGAACACACCACTCTCAATCCCGTAACCTTCACTTTCCTTTTCTCTTCTCTATTATTTTTT TTAATCATTGAAAGATAGAATAATATCTATAATTTTATTCAAGTTTAATTATA
>Gm03_3	TCCTCTCCTCACCAATTTGTTTTCCATATGAACAAATTTATTACACAGGATCAAGGTACTCAC

8154653	ATCTCTTAATTCTGTATTATATTGGTTCACTTAATTTTATGAACCTTTAAAATAAATGCAAAAG GCACTTTCTTCTGTAGATGGTTTCTTTTCTTTCTTTCCACCATCCTGGACAT[G/T]CGTGGGA ACAAAAATGGAACATACAGCAGCTGCATCTGTTTGAGCTTAACTCATTGACCATAAGTGAGG CAGAAGTTTGGTAGTTTATGTGTTTACCTATGCTTTCTTCTGCTCATGCTGATTCCGATACTC TCTCAACAGAAGTTCAGCGTAGAGAATTTCACTCCATAAATCAGGAAC
>Gm03_3 8170431	TTAGATATCTTAGTATGGGAAAATAAAACGAAAATGTGAAATCTTTCAGCTGTGGAAAGAT GGAGAGATGAAAGAAGAAGTCATTGGAGGCCATAAGGCCTGGCTAGTTATTGAGGAAGTC AAAGAAATGATCCAAAAGTATTTATGATTCTTCTATTTTATTTTCTTGCTCCCTCA[A/C]CCC CCCACCCACCTCACAAAAAAGAAAGAAAGAAAGAAAGTTACAATTGAAGAAACTAA TCTTCCAATGTTTTACTTTTTTCTTTAACATTTTTCATGTTGGCAACCGCATGTAGCTGTG TATCATATATATATTAATGGAACGTGTGTACCATAGCTGAAAGAAATTTA
>Gm03_3 8170524	ATAAGGCCTGGCTAGTTATTGAGGAAGTCAAAGAAATGATCCAAAAGTATTTATGATTCTTC TATTTTATTTTCTTGCTCCCTCAACCCCCACCCACCTCACAAAAAAGAAAGAA AGAAAGAAGTTACAATTGAAGAACTAATCTTCCAATGTTTTTACTTTTTTCTT[A/G]TAACA TTTTTCATGTTGGCAACCGCATGTAGCTGTGTATCATATATATATTAATGGAACGTGTGTAC CATAGCTGAAAGAAATTTAAGTGAAAAGGAAGATGCTCACTGTTTCTCCCTCCCTCTCAAC CATTAGAGATTTAGAGGGGAAAAAAGTGGGGGTTTTATACCAAATAA
>Gm03_3 8173525	ACCTCCTTACTAATCCTCGCATTGCTCTGTGTAACCCCTTCCACCTATTCTCATCCATCAAA ATCTCAAACAAATCTTAGCTCTGCAATTAACCTCTCCGTCATGCTAATACTCTTCCCTCCC TTCTTCACCTTCTCCTTCTTCTCCTCCTTCTTCGGCGCAGCAGCCGTTGTTTT[T/C]GTCGTC GTCGGAGTCGTCGTTGTCGTTGTCGCCGCCGAGACGGCGACAACGACGACGGCGCAAC CTTCTTGGGTTCCAATTCGTCCTTAACAGGACCTCCTTTGGCCATGCTCTGCACCCAAACC CTAACCTTCTCCAAGATCAAGGGCTTCCCCTTGGAACCATGGCATCCTTAATC
>Gm03_3 8173719	CGTCGTTGTCGTTGTCGCCGCCGAGACGGCGACAACGACGACGGCGCAACCTTCTTG GTTCCAATTGTCCTTAACAGGACCTCCTTTGGCCATGCTCTGCACCCAAACCTAACCTTC TCCAAGATCAAGGGCTTCCCCTTGGAACCATGGCATCCTTAATCCTCTTTTCCAACCG[G/T] TCCCTCGTCGTCGACAGTAACCCCTAACCTCGGGATCCTCGTCGGCGTTCTCGTCGGAGAT GTAGGGGATCTCGACGGTGCCGTCGACCTTGAGAAGAGAGGTTCCAGGGTGTCTTTGGC CTCTCCCTGCCAATTGAGAGTGAGGCTGATCTCGTAGCCGGGGATGATCTTCCCCTTTTCG
>Gm03_3 8173815	GCTCTGCACCCAAACCCTAACCTTCTCCAAGATCAAGGGCTTCCCCTTGGAACCATGGCA TCCTTAATCCTCTTTCCAACCGGTCCCTCGTCGTCGACAGTAACCTAACCTCGGGATCCT CGTCGGCGTTCTCGTCGGAGATGTAGGGGATCTCGACGGTGCCGTCGACCTTGAGAA[A/G] AGAGGTTCCAGGGTGTCTTTGGCCTCTCCCTGCCAATTGAGAGTGAGGCTGATCTCGTAG CCGGGGATGATCTTCCCCTTTCCGACATTGATGTAGGCCTCGCCGTCGAGGGAGCGAAGC GACGTCGTTTTGAGGAAGAGGTTACCCTCGCCGTCGAGGATAGGGAGGTTAGTAGAGGAG
>Gm03_3 8174150	CGAGGATAGGGAGGTTAGTGAGGAGGCTGGTGAAGAAGGTTTGACCACTGAGGCAGTT GGTCTCGGACCAATGCCAGTTGTGGACATTGGTGCCGTCGGGGCGTCCCTCCACGATCCA CCGCTTGTGCCCTCACCGTAACGAGCCATTGCAACTGATGCCTCTTTCTGGGAGAGT[A/ C]CCTTAGTGATTAATAATAATTGGATTGGGGTGAGGGTTATGAACTATGATGATTATTATTA TATGTGTTGAGTGTGTGTGGAATGGTCTAGAATGAAAAGGAAGGGTCTAGAATGTTGATGT TGCTTTGGGTACACTTTCTTACCAACCAACCAAAGCCCGAGAAGGTTCCCTAGGGTT
>Gm03_3 8389430	ATAAACACCACACTAAAATTATGCCCAAACACACAAATAAAAAGAGAATAGATATCTCGATT CACGAGAAACTCAAATAAATGAAACTATGCAGGAACCATGTTAGAAAACAACCTAATTATTT TCAAATGTTAAAGGACTATACATCATCCAATGTCAATCCTAATCTTACTATATAA[T/G]GCAAC CAACTACATACGAAAACAAGAGCCTTAAATCTAAAACCTCCTGACTAGCAGATCCAATCTCGG TCTTCCCCTTGCCAGCGACCTTCTTGGGCAACAAATTCTGGTGAATGTTAGGCAGAACACC ACCGTTGGCAATGGTCACAGATCCCATCAGCTTGCTCAACTCCTCGTCGTTT
>Gm03_3 8389449	TATGCCCAAACACACAAATAAAAAGAGAATAGATATCTCGATTACGAGAAACTCAAATAAA TGAAAACCTATGCAGGAACCATGTTAGAAAACAACCTAATTATTTCAAATGTTAAAGGACTATA CATCATCCAATGTCAATCCTAATCTTACTATATAAGGCAACCAACTACATACGA[T/A]AACAA GAGCCTTAAATCTAAAACCTCCTGACTAGCAGATCCAATCTCGGTCTTCCCCTTGCCAGCGA CCTTCTTGGGCAACAAATTCTGGTGAATGTTAGGCAGAACACCACCGTTGGCAATGGTCAC AGATCCCATCAGCTTGCTCAACTCCTCGTCGTTTCTTAACCGCCAGTTGAATGT
>Gm03_3 8389457	AACACACAAATAAAAAGAGAATAGATATCTCGATTACGAGAAACTCAAATAAATGAAACT ATGCAGGAACCATGTTAGAAAACAACCTAATTATTTCAAATGTTAAAGGACTATACATCATCC AATGTCAATCCTAATCTTACTATATAAGGCAACCAACTACATACGAAAACAAGA[G/A]CCTTA AATCTAAAACCTCCTGACTAGCAGATCCAATCTCGGTCTTCCCCTTGCCAGCGACCTTCTTG

	GCAACAAATTCTGGTGAATGTTAGGCAGAACACCACCGTTGGCAATGGTCACAGATCCCAT CAGCTTGCTCAACTCCTCGTCGTTCCCTAACCGCCAGTTGAATGTGCCTCGGA
>Gm03_3 8419312	TGCTTGCAACCCCTTTTGGACCAAGGGGAGAAGCATTGAAGGACTCAAATTCCTAAAAGT CTTGTTTGTGTGGCTGGTTTGGATCTTCTGCAGGATTGGCAATTGGCATATGCAAAAGGTCT TGAGGATTGTGGACAACAGGTCAAGCTTCTTTTCTCAAGGAAGCCACTATTGGCT[G/C]CT ACTTCTTGCCAAACAATGACCATTTCTATTGCCTCATGAAAGAAATCAACAACCTTTGTCAATT CTGATTCTGACTGTTAATATACTATTAGTACTAAGACGACAGACGACTACTACTACTACTGC TACTACTTAACCTTACCTTACCTACAGGAGGCATACATAACTAAAGCTTGACA
>Gm03_3 8419319	AACCCTTTTGGACCAAGGGGAGAAGCATTGAAGGACTCAAATTCCTAAAAGTCTTGTTT GTGTGGCTGGTTTGGATCTTCTGCAGGATTGGCAATTGGCATATGCAAAAGGTCTTGAGGA TTGTGGACAACAGGTCAAGCTTCTTTTCTCAAGGAAGCCACTATTGGCTTCTACTT[A/G]TT GCCAAACAATGACCATTTCTATTGCCTCATGAAAGAAATCAACAACCTTTGTCAATTCTGATT TGACTGTTAATATACTATTAGTACTAAGACGACAGACGACTACTACTACTACTGCTACTACTT AACCTTACCTTACCTACAGGAGGCATACATAACTAAAGCTTGACATTTTACT
>Gm03_3 8419321	CCCTTTTGGACCAAGGGGAGAAGCATTGAAGGACTCAAATTCCTAAAAGTCTTGTTTGT GTGGCTGGTTTGGATCTTCTGCAGGATTGGCAATTGGCATATGCAAAAGGTCTTGAGGATT GTGGACAACAGGTCAAGCTTCTTTTCTCAAGGAAGCCACTATTGGCTTCTACTTCT[C/T]GC CAAACAATGACCATTTCTATTGCCTCATGAAAGAAATCAACAACCTTTGTCAATTCTGATTCTG ACTGTTAATATACTATTAGTACTAAGACGACAGACGACTACTACTACTACTGCTACTACTTAA CCTTACCTTACCTACAGGAGGCATACATAACTAAAGCTTGACATTTTACTTG
>Gm03_3 8717673	AAAAATAGAAACAATCAAATTACCTATACATATTTTTTGAATTGGAATTTATTTAATTCCATTT GCAAAATTATTGACCTTGAGATTAGAACATTTGGCATCCTGAAGCAAGAGAGTGAAGTGA AACCGAATCTATTTGAGTCAAGATCAAACCTGCATTAAATTATCCTCCAACCTGAT[T/C]CCCTC CGATTACAATGGGAGTTCTCCCGCGGGTCCCACCGTCCACAAAACCCAAACACACACATC CACTCCTCCCTTCGCAACCCCTACCATCGAATTCCCGGCCAAAGATTCTCCAGAACACGTCC TCGCTGTGCATCACCAGATCAACGGTGGGCACCGCGGGTCCAACCTCGTGTCTC
>Gm03_3 8717682	AACAATCAAATTACCTATACATATTTTTTGAATTGGAATTTATTTAATTCCATTTGCAAAATTA TTGACCTTGAGATTAGAACATTTGGCATCCTGAAGCAAGAGAGTGAAGTGAACCGAATC TATTTGAGTCAAGATCAAACCTGCATTAAATTATCCTCCAACCTGATGCCCTCCGA[G/A]TACAA TGGGAGTTCTCCCGCGGGTCCCACCGTCCACAAAACCCAAACACACATCCACTCCTC CCTTCGCAACCCCTCACCATCGAATTCCCGGCCAAAGATTCTCCAGAACACGTCTCGCTGTG CATCACCAGATCAACGGTGGGCACCGCGGGTCCAACCTCGTGTCTCCGTGAGATC
>Gm03_3 8718672	GACGGGGTTTTCGGGGAAGAGGGGCGCAGAGGGAGGAGTTGTTGGAGCATGCGTTGGAGG GGAAGGAGTTGCAGAGAGGGGTGTTGCAGGGGATGTGGTGGGAGGAGGAGGAGGTGTAG GTGGAGTCGCAGAGGAGCCAGCTGAGGGAGGAGCCAAGGTGGAGGTGGAGTTTTGTGGG TT[G/T]GAGTGGGGTTTTCAGGAACACGGAGAGAGTGTAGAGTTGGGTGGTGTGCTCTTTT GAGATTGGGGCTATGAGGGGGATTTGAAAAGTGGGTGTGAGGAAGAGGAAGAGTATCGCC AAGTTGAAGTTGCAGAGAGATGGAAGAGGAGGAGGCATTTTGGTTTGCTTGCTGGAGCAA GGAGA
>Gm03_3 8718710	TGTTGGAGCATGCGTTGGAGGGGAAGGAGTTGCAGAGAGGGGTGTTGCAGGGGATGTGG TGGGAGGAGGAGGAGGTGTAGGTGGAGTCGCAGAGGACCCAGCTGAGGGAGGAGCCAA GGTGGAGGTGGAGTTTTGTGGGTTGGAGTGGGGTTTTCAGGAACACGGAGAGAGTGTAGA GT[T/C]GGGTGGTGTCTGCTTTTTGAGATTGGGGCTATGAGGGGGATTTGAAAAGTGGGTGT GAGGAAGAGGAAGAGTATCGCCAAGTTGAAGTTGCAGAGAGATGGAAGAGGAGGAGGCAT TTTGGTTTGCTTGCTGGAGCAAGGAGATAAACAAGAAACAGTGTTGGTGTGGAGTTAA AGTG
>Gm03_3 8720114	AACCTAAGCTTCTTTTCTATGTGATGCATGAGCCGAACAATGCCGTAGCACAAATACAATCT CCTAGAAAAAATTGTACAGCTTTTCTGCTGGCACTGCATTACAGGAGTCTTTGCTAGCTCTTC AATAAAAATACGATGTCGTTTCGACATGCTTAGCTTTTATTATAACTCACTTGCGC[G/A]GCGA AGAGATCATATGCTGGTGAGCCTTGATATCCCTTTGTATTCTGTCTGCATACAAGGGTAGGC GAATGATCTGATTGATCTCATTCAATAAGACCTCTTCTGTGATGTTATCCTTTATGCTTCGCA TTACCTGTGTTAAAAAATTAACAAAACTGTTTCAAGAGGCTTTTACATGA
>Gm03_3 8720115	ACCTAAGCTTCTTTTCTATGTGATGCATGAGCCGAACAATGCCGTAGCACAAATACAATCTC CTAGAAAAAATTGTACAGCTTTTCTGCTGGCACTGCATTACAGGAGTCTTTGCTAGCTCTTCA ATAAAAATACGATGTCGTTTCGACATGCTTAGCTTTTATTATAACTCACTTGCGCC[G/A]CGAA GAGATCATATGCTGGTGAGCCTTGATATCCCTTTGTATTCTGTCTGCATACAAGGGTAGGC GAATGATCTGATTGATCTCATTCAATAAGACCTCTTCTGTGATGTTATCCTTTATGCTTCGCA

	TTACCTGTGTTAAAAAATTAACAAAACTGTTTCAGAAGGCTTTTCACATGAG
>Gm03_3 8942898	ACGGAGTTCAATTTACTGAATCTAATTTCTTTTTTTTTTCGCATTTCTATTTTTCTCCTCATTT CCTTGGAACCAAACAGTTAGATTAATGACAGTTAGATTGATGAGGTACCTGAATCCTGACA TAGTCGGTACGATTCCAACAGAAGGCGTTCCCTAGCATCTGGTCAACGGTCT[A/C]TGA AAC ACCGTCGAGCGCAATGTCAATCAGCATTGACGTAGAGCACTCGCCGGCATTGTTTCATTCTC TTCGCCGGAGCTCCGTTTCCGATGGAGAGAACAAGGAGATCCTCCACGCCGTTACCGGAC GGGAAGTCGCGCTTGTTGTGGAGGACGTGCGTGACCGCCGCCGCGGCCGATT
>Gm03_3 8942920	TAATTTCTTTTTTTTTTCGCATTTCTATTTTTCTCCTCATTTCTTGGAACCAAACAGTTAG ATTAATGACAGTTAGATTGATGAGGTACCTGAATCCTGACATAGTCGGTACGATTCCAACAG AAGGCGTTCCCTAGCATCTGGTCAACGGTCTCTGAAACACCGTCGAGCGCAAT[A/G]TCAAT CAGCATTGACGTAGAGCACTCGCCGGCATTGTTTCATTCTCTTCGCCGGAGCTCCGTTTCCG ATGGAGAGAACAAGGAGATCCTCCACGCCGTTACCGACGGGAAGTCGCGCTTGTTGTGG AGGACGTGCGTGACCGCCGCCGCGGCCGATTGTTTCATCACTAAGCCGCCGCTCG
>Gm03_3 8942927	TTTTTTTTTTCGCATTTCTATTTTTCTCCTCATTTCTTGGAACCAAACAGTTAGATTAATG ACAGTTAGATTGATGAGGTACCTGAATCCTGACATAGTCGGTACGATTCCAACAGAAGGCG TTCCCTAGCATCTGGTCAACGGTCTCTGAAACACCGTCGAGCGCAAT[G/T]GCAT TGACGTAGAGCACTCGCCGGCATTGTTTCATTCTCTTCGCCGGAGCTCCGTTTCCGATGGAG AGAACAAGGAGATCCTCCACGCCGTTACCGACGGGAAGTCGCGCTTGTTGTGGAGGACG TGCGTGACCGCCGCCGCGGCCGATTGTTTCATCACTAAGCCGCCGTCGACGGCGG
>Gm03_3 9790817	CGTGTGTTGCCATGCAGGGTTTTGGAATTTTGGCAGGTGGTATCTTTCGAATTATAATTTTCAG TTGCATTCAAGGAAAGGTTTGATGCTCCACCATATGAGCTTGATCCAGCTGGCTCAACTGTT GCACAAGCAGACTACATTTGGAGGATAATTGTTATGGTGGGAGCACTGCCAGCTG[G/T]GTT AACTTACTACTGGAGGATGAAGATGCCGGA AACTGCCCGTTACACCGCTCTAGTCGCCAAG AACACGAAGCAGGCTGCAGCAGATATGTCTAAGGTTCTGCAGGTTGAGATTCAAGCTGAAC CGCAGAAAGAGGAGCAGAAGGCTAACTCATATGGCTTATTTTCAAAGGAGTTCTCT
>Gm03_3 9790874	TTCAAGTTGCATTCAAGGAAAGGTTTGATGCTCCACCATATGAGCTTGATCCAGCTGGCTCA ACTGTTGCACAAGCAGACTACATTTGGAGGATAATTGTTATGGTGGGAGCACTGCCAGCTG CGTTAACTTACTACTGGAGGATGAAGATGCCGGA AACTGCCCGTTACACCGCTCTAG[T/C]C GCCAAGAACACGAAGCAGGCTGCAGCAGATATGTCTAAGGTTCTGCAGGTTGAGATTCAAG CTGAACCGCAGAAAGAGGAGCAGAAGGCTAACTCATATGGCTTATTTTCAAAGGAGTTCTCT CCGTCGCCATGGACTGCATCTACTTGGTACAGCAAGCACATGGTTCTTGCTTGATAT
>Gm03_3 9794956	AAGTTTTGAAATTTTCTATAGATACTTGCCAAAGTTAGAACAAGGAACCTTTTCATCCAAGTA ATCTATCTGGATCCATAGAGTCACGCTCATACAATACAAGTCCAGCACACTGGGATTTACAA CAATAACTGCCAAAAGCTTTAAACCCATCAGCCTTCAGATTCAATATTATTGGAT[A/G]CGAGT TATATATCTGTGGCAAGATGGGCTCTGCAGATGACAAAGCATGAAGAGGTAATTCATTAAA GGTAAACCTAGTAGATTCTGATATGAAAATGAATTTAAGTGCTTAAACATTTCAATGTGCAC GGTCTCTCAGCACTTCAAGATCCATTTCTGTGGGGTCAGTAGCCTGGATT
>Gm03_3 9795164	TCTGCAGATGACAAAGCATGAAGAGGTAATTCATTAAGGTAAACCTAGTAGATTCTGATA TGAAATGAATTTAAGTGCTTAAACATTTCAATGTGCACGGTCTCTCAGCACTTCAAGATCC ATTTCTGTGGGGTCAGTAGCCTGGATTTCTAATGGACACCATCTAGCTCACGTC[A/G]AAA CCTGTCTTTAGTAGTGGCATAGAGCATCTTGGCACGGATCCGAGATGTTGAAGGGGACCTA ATCAATATCATGCAATGAATACTAAGAGTGATATCTCTGGCAAAAGCATAGAATGAGTAAG TTAAGAAGCATGTTGTTGAGGATACATGAGAAGGCTGGGATAAAAGAAGTTCAA
>Gm03_3 9795203	GTAAACCTAGTAGATTCTGATATGAAAATGAATTTAAGTGCTTAAACATTTCAATGTGCACG GTCTCTCAGCACTTCAAGATCCATTTCTGTGGGGTCAGTAGCCTGGATTTCTAATGGACAC CATCTAGCTCACGTCTAAACCTGTCTTTAGTAGTGGCATAGAGCATCTTGGCAC[C/A]GATC CGAGATGTTGAAGGGGACCTAATCAATATCATGCAATGAATACTAAGAGTGATATCTCTGG CAAAAGCATAGAATGAGTAAGTTAAGAAGCATGTTGTTGAGGATACATGAGAAGGCTGGGA TAAAGAAGTTCAATCCAAGTTTGTCTTCAACAAGAACAAGAGAGGAAGACAA
>Gm03_3 9846797	AAACCTCAAGTTTCAGGTAAAAACAGTGACATGCAATCCTCCTTTACAACAAGCAAGTGACA AAACACAACTAAACAAACAAAGTAGTCATTTCTATATACTTACAAAACAAACAAATTATTAT CATAAATAAACTCAACAAGAGTCCGATTTTCTAATCTGTACTCCAATGTGCG[T/C]GATTA GTTGTCTCCACACTCTATGCACTAACAACAGTACTAACTTCATCCCCTTCTCGCATTAATAAT ATAAGGAAGCGGATCAAAATGTGGGAACCACTTAATATTATGAGAACTAGCATGATCTCAC AAGGATGTTATCACAATCCTCGGGATGGAACTCCTGCCAATTACACATT
>Gm03_3 9963852	AGAAATAAGCTAAACATGCATCAAAATAAATATACAAGAAAAACAGAGCTAGAGAGGGTT CAATTGACCTTGAAAACGATCCATCGAAGAGCTCTAACGCGGCGGAACGGTGAGCATCGG

	AGACGGGGGCCGGAATCGTGACGGCTCCGCAGATCGAAACAGCGAAAAGTAGCGCCA[G/T J]TAATCCTCCTAGGTTTCGGGCCATTCTGCACCGATCAATGGAAGAATTTCAACAACAAAG CGAATCGCATTTACAATTTTCGATGCGGATCGGCTTATGTTAGGGTTTGAGATCGAGAAGAA GAGAATTAGATCATACGCGGAAGCAGCAAGAAGCTGAGAAGAAATGCAATTGGCAAGC
>Gm03_3 9966061	ATTTTGAGAATATTACATTGACTGCAAACACAACCTCAAAATCCATTAGTGCAAATTAACATTC GTCGCTTCATTCTAATAACGTGAGCCAGATTTTGATGTAACCTGGTTCACAACACAAAAGGCA TAAATGGCTCAATCACTTGACACCAGGCACCAGCACTAGTTTCAAAAATACTGT[T/C]TACG CATGGAACAAAAGACACTTAAAAATTTAGAGCTTCTACATCACTCAATACTTTGGAGGTGGCT GCTGGGTTGGTAATAATCCAAAACGCTTCTTCAGAAGAACTCTCTGTCTTGAATATTTATCA TCGGGTGAAAACCGGGCTGCAGTTCAGGAAAACCGGTAAGTACAGAAAATG
>Gm03_3 9986148	TTGTGAAGATGCTGCATGCAATCATTTGTTGAAATTTTTATGGCTGGAGTTTTATTATACTT TCTACTCGTGTGTTCTTAACTTTTTATTTTTATCTTGTCTCTTAAGTATCATGATCAAGAGAA GGGACGCAATTCGTCCCGATATGGCAGCAGTTCAAGAAGACCCATAATCTCA[G/T]CGTCAA CCAGGCCAAGTTCCTCTGGTGATCATACTGACAGTCGTACTGGCCGGCTAACCTCAAGTGG AAGCCGACAATCTGCCACACATAGAAATATTCAACCTATGCACGAGACAAAACAACCAACTT ACACACGCTCTGGATCCACCAGAGGCAACCGTGATGATCCTCTGCGGAGTT
>Gm03_4 0013399	TTGATATGAAACCAAGTACTGTACAGTTAATTCATGACACGAACAAATGAAGACAACCTTGG TGGGTTGGGGGAAAATATTATTGTAACATTTCAATGTAGTAACTAAGCTGATTAGTTGTTGA TACTTATTCGTGAAACAGGTTTTGTTCTTGCGAGACCAACTCCATTTTCTCATGA[A/G]TGAT ATCAGTGCTCCTTCTCCCTCATGATACTATTGTCTATCAGTTTTGACTAATGTAACGGGTTT CTACAATACGATGACATTAGACTGTTTTAGCATCAGGTGTGTCCTTGAACCTTGGAAATCAATA CAGCATTAGCATTGCTGTGGTGCTATGTTGAACACGTCAAAAAACAACCTACA
>Gm03_4 0013405	TGAAACCAAGTACTGTACAGTTAATTCATGACACGAACAAATGAAGACAACCTTGGTGGGTT GGGGGAAAATATTATTGTAACATTTCAATGTAGTAACTAAGCTGATTAGTTGTTGATACTTAT TCGTGAAACAGGTTTTGTTCTTGCGAGACCAACTCCATTTTCTCATGATA[T/C]CAGTG CTCCTTCTCCCTCATGATACTATTGTCTATCAGTTTTGACTAATGTAACGGGTTCTTACAATA CGATGACATTAGACTGTTTTAGCATCAGGTGTGTCCTTGAACCTTGGAAATCAATACAGCATT GCATTGCTGTGGTGCTATGTTGAACACGTCAAAAAACAACCTACAGTTGTA
>Gm03_4 0021724	GAGTTTCTTGACACTGTGACAATTGTTGGGGGTCCAAAGTCAAGGAATGAAGTGTGGACAG TGTTTGCCACTGCACATCAGATGACAAGTTGTAGAATGGTTGCCACTGAAGAAGAAGGTGC ACTTGATGAGAATGGTGAGAGTTGGGAAGCAAAAGGCTTGTATGTGTGTGATGGCAG[C/T] GTTTTGCCAAGTGCAGTTGGTGTCAACCCCATGGTAACCATTCAGTCTACTTCTTACTGTAT TGCCACTAAGATTGCAGAATCACTTAGCAAACCATGAAAAATCAAGTTGACTGTTTCACGA AAAGAAAAAATACTATGCATTCTTACACTGTCATCCATTAGAAATATTATTTAT
>Gm03_4 0059714	ATTGATATTCATTGTTAAAAACTTGGCTAAATCTCTTCTTTTAAAGTTCTACACTAATATTTCCA ATTGTTGGTAGCAGGAGCCAAAGCTATTCAATGGCTTAAAGTTTTATTTTCTGGGGATT ATTGTGTCAACTTATAAGGAAGATCTCGAAGAAATTAATTGAAGTGGGAGGAGGT[A/C]CTGTT TTGAGAAGCAAGGAAGAATTGGAATCACAAGACATGAATGCAAAGGAGATTCTTCACAGT TGCTGATTGTTTACAACCTTGATCCTCCTCAAGGATGCAAATTGGGGGAGGAAGTTTCAATT CTTTGGCAAAGGCTAAACGATGCAGAGGATTTAGCTGCTAACACTTTACAAG
>Gm03_4 0084046	CAACAAAGCATGTTTGATTTTTGGGATTGAAGTTTATTTATGATTCTGATAAAATGGGTTTGG GATCTAGGGCTAAACCTTTGGAGCAGATAAACGTGAGCCCTAAAAAGAAAGTAACTTTTGAT GTCAATGTGAAAACGTACGAGCCCGAGCCTGATGAAGTTGCTGATTACCAACTT[G/A]TGAG AAGTGAAGAGGGTAGAGAGGAAAGTGCTTTTGTGAAAAGTTAAGCCAAACAAAGTCTTAC TCTGAAGTAAGTTCGGTGACTTCAGCGAGGTCTTACCCTACAAACCATAGGTACCACAGTT GCACGTGTAGTGATGATGAAGATGGAGCAATGGAGTATTGGGATAGTGATGTTA
>Gm03_4 0084245	AGAGGAAAGTGCTTTTGTGAAAAGTTAAGCCAAACAAAGTCTTACTCTGAAGTAAGTTCCG TGACTTCAGCGAGGTCTTACCCTACAAACCATAGGTACCACAGTTGCACGTGTAGTGATGA TGAAGATGGAGCAATGGAGTATTGGGATAGTGATGTTACTGATGAGGATGAGGATG[A/G]T GATGATGATGGTGATAGTGACATGGGAGAAGAGTATGATGAAGTGGAAGAGGATTTTGAAG ATGGAATAGTGTACTCAAGGTCAAGAAATGGTGCTAATCAAGTGGTTGTTGGAGAGGTGGA AAGTCCAATTCCAATGCATGATAAGGATCTGAAGTCAATTGGGTTGAACCTCAATGT
>Gm03_4 0132046	GATTCATATGACCCAGCATTATCTGCCAAAACGCAACAATAACAATTAGAAAACAGAAA AAAAAATTAACAAAGACAGAGATCAATAATAAGGAAAACAGAAGAAAAAGTGAAATACCTTCG AAGTACATGTAAACACCGTCCCTTTCGGCGCCACGGCTTGCGCTGACGGACGATGA[A/T]GG CGGGGAGCACCTTTTTACGGAGATCAGGCTTCCCTTCTTGACGGTGGCCATAACCATGTC ACCCACGCAAGCTGACGGCAACCGGTTACGCTTCCCTTGATCCCTTCACTGATATGATG



	TAGAGATTCTTCGCGCCGGTGTTCGCGCGCAGTTCACCGTCGCGGCCACCGGCAG
>Gm03_4 0154304	TTATTCTAATGTTGGTGACGTGAATTCGTATGCTGGGGATCCGAATTCATCCTACAGCAAG CTCAATTCAGTGCAACTGGTGAGTCCAAGCCAGCTGGTGGAGCGGCTGATTCTAACGAGG CCTCTGCTGGGGTTGGAAGCACAGCTGCTGACTCTACTATGGTATCTGACTACAATT[C/T]A TCTGTAAATGGTGGTGTTCGCGGTGCAGTGACAAATACTTCTGGACTTGAAAACGGGAATG CGTTGGAGAATGCTGATGGGTGCGCTGATGAGAAACAGCAAGCAGATGGTTATGGTGTGC CTTTTTCTTTTACCTTTTTACTTTCAATGGAATTATCATGTTTTAAATTTTATTTTTG
>Gm03_4 0154315	TTGGTGACGTGAATTCGTATGCTGGGGATCCGAATTCATCCTACAGCAAGCTCAATTCAG TGCAACTGGTGAGTCCAAGCCAGCTGGTGGAGCGGCTGATTCTAACGAGGCCTCTGCTGG GGTTGGAAGCACAGCTGCTGACTCTACTATGGTATCTGACTACAATTCATCTGTAAAT[A/C] GTGGTGTTCGCGGTGCAGTGACAAATACTTCTGGACTTGAAAACGGGAATGCGTTGGAGAA TGCTGATGGGTGCGCTGATGAGAAACAGCAAGCAGATGGTTATGGTGTGCCTTTTTCTTTT ACCTTTTTACTTTCAATGGAATTATCATGTTTTAAATTTTATTTTTGGACAATAAATT
>Gm03_4 0172340	CAGACTACGGTTTGGAGGAAGAAGCCAGGGAATATGTCAACTCAGTTCATATAGATGATGA TCCAGTGGACGAGTATAGTCTTCCTGAGCACCAGCAGCAGCTACAAGAAGAACTTGAAACC GAAATTGTGGAGGAGGAAACTCCCGTACAGAAAGGAGGCATCTCCACCAATTCATAGCATT[A/G]C ACACACTGTCCAAGAACCCCTGTTGCTCTTGTGGAAGAGTCCTTTGAGGAGCCTCCTAAG AAAACATATGCATCTATTGTATGTAATACTTTCTATACATGTTCTTTTTTTTTTGGCAGACTAC CTTTTTTGTGTGTAGTTATGTACTATCCATAAAAAAGGTACCACATGACTGGA
>Gm03_4 0172927	CTGTATGTTGGACTTAAACTAAATAAAGAGCATTTATCTTTTTTAACTTTTGGTGTGTTGTCT TAATTTAATCGTGTTCATGAACCTGCAGTTACGAGTTTCAAAGGGCAGCCAGTGTGTC AGCTGCTCCACAGTATGCTCCACAACATTCTTTTAAAAGTGCCCCACCTCCTT[A/G]AGAGTT GAACCATGTAGCACAGCCTGCTGTTTCAGCAGTCAAGCTCTGCATCTATGTATGTTCCCTGAG TCAGGGATTGAGGCCGAGAAAGAGGCTATGGACTAGAAGAAGAAGGTTAGTTTATGTTT CTGATATTATAACCTTTTCTTGGCATATTTGGCGTCTTCATGTTTTTGAAT
>Gm03_4 0179701	CCACCATTGAGCAGCACTGAACCCATCTTCACTCTCACTTTCCATTTCTTCATCTTCTCTTCT TCACTCAAGCCATTGTGCTTGGCAGTAGAGAAAGAAGGAATGGTCTCCAGCACCAGCTCC TGTGTCTAGCTGAAAACCTCTTGGAAGGGTCTGAAAACCTCAAACCTTTCTTCATT[A/C]TTGC AAGAAAAGAAGGCCAAGTCTTGCAAGTTCCCATCAGATTGGACTCTTCTGATCAAGCCTTT GTTTTGCCTTTCAAGCTCACCAATTGAGGGAGAGATTGGAGAAGAACCACATGCAGAGGAC CCAGTTTGGTGGAAGTAAGTTTGTGATAATGTTGTGGGACAGTGGTTGGTGA
>Gm03_4 0211426	TGGCGTACCAGCCGGTGTTCGGAGAAGACGCCGTTGACGTTTTCCAACGGAGGACCGAGG CCGGCGTTGGTAGTGAACCTGCACATGTTGGTCCAAAGGGATAAGCTTCTGCAATGCTTCA ACATGTCCTCGTTGAAGCGCGAGGGAAGGTCTGGACGAAGATGTAGCGTCTCCGCAT[G /A]GGTCGCTCTTGTGTCGGCGGTTTCGGAGGGCACGGTCTGAAGGGGAAAGTGGGTTCTT CGGTTTCGGAAACTCGCTCAGGGAAGCCGAGGTTTTTACGAGGTTGTTGGTGTGTTGTTGG GATTGGAAAGTGATGGGGACTGCTGGGTGAGGGGCGGTGGGTTGGTGGCGGAGGACGGC GA
>Gm03_4 0288569	TCATCATCATGTTAATGCTCAAACCTGTGCTGTAGTTTTGTAAATCTCTTTAATTTTATTGTT TTCTTTTCAAATCTTTCAGGACATTCAGGTTGCATTGCGTGATGAAATCAAAAGTGCTTGTG AATATCGTCCATTTGAGAAATGTGATTATGGTTCAAGAATAGCCAATGAGATT[A/C]GTCACA AGAAAGTGGAATTTATTATGTCCAGTTGGATGCTATAAAACAACTGTAGATGAGTATCAT GTAGCAGTTTGAAATGTAGCTTAGTTCCTGTAGAAGCCTGGGTTACCTCCCGGTAATTG ACTTATTTGGGATTGCTTATTACTAGTTAGCATGTTGTTTTGGATGGGGT
>Gm03_4 0338539	CATGCTTACTTCATACTAAAGATTATACAGAATACAAAGAGTGTTAGATGTGAGAGCAATT CACATCAGAAAATTTAAGATTGAGAGAGAGAGAGAGCATTATTTTACCTCTAACACTGTA TATATAGGCAATGTTGTTTCTCCATCGATAACAATACTTTTAAAGAAGAGAACCAT[G/A]ACGC CGACCATACGCAAGTAGTATGTGCTCAGATGTAGGCGAGAACTATATCAAACAGCAATTCA ATCCTCTAAGAGAAGGCCATTTCAAATTATATTTGACCAATTTAATAAACTTGGCCCTAAA AGTTGCAACATGAGTTACTTTGGTTGCTGAAAGTGCAATGCTATACAAATCA
>Gm03_4 0339831	TAACATCAGAAGGTACCTGCAGAGGACAGCATGTGGTATGGTTAGACGACATCTATCAGCA TTAAGTGGGGCACAGGGATTTTTTATGTCATGTGACCATACTCTTAACCTTCACAGTACAAGG CAACTCTGCAGCTGTTGCAGCCACTGATGTGGCAAGTTCTGACTGAATTCGACTTA[C/T]AA TGGTTTGGATATCGAATCCATCATGTGGGGCATTAAAGAAGCAGCAAGATTGCCAGATT AGATACAGGTACACGGGACTGTGAAAAATGATTCCGTAAACCTGATCTCATTGAAGAACCC GGTATGCTGATGCTGCCAGGATTGCTGAAGATGGCATCGCTGCATCAACATTAGA
>Gm03_4	TGTTGGCTTCCTTCAGCAGGTCTGGTTTCATCAGTCTCCATACCATTGAGATTACT

0340465	GAGGCCAATTCCCATTCCATTAGAGAAAGCAGTATGAGCAGGATATTCTGCACCAGTTTGA GAGCTAGAGGGCATCTCAGAGAATGTATCCATGGGAGACACGGTAGTCTCATAATG[C/G]G CTGCTGCACTTGGGTCTGCATGTAACGTACCATGGCAGCAGACTCAATTTGCATGCTACC TGAGCCAACATCATGACTAGCATGCTGTAGTTCTGCTCTTGACTCATCAACAGTATATGCCG GCATAACATAGAAAGGTAATGACACATAAGGTAGTTCGCTGGACAACTAATATGT
>Gm03_4 0340517	TGAGATTACTGAGGCCAATTCCCATTCCATTAGAGAAAGCAGTATGAGCAGGATATTCTGC ACCAGTTTGAGAGCTAGAGGGCATCTCAGAGAATGTATCCATGGGAGACACGGTAGTCTCA TAATGCGCTGCTGCACTTGGGTCTGCATGTAACGTACCATGGCAGCAGACTCAATT[G/T]G CATGCTACCTGAGCCAACATCATGACTAGCATGCTGTAGTTCTGCTCTTGACTCATCAACAG TATATGCCGGCATAACATAGAAAGGTAATGACACATAAGGTAGTTCGCTGGACAACTAATA TGTTCTGTAGGATGAATATTTGTGACAAAAACAGCAGGCGGAGGATATTGTAAT
>Gm03_4 0377254	AAGTTGCCGCAAGAGAAAAATACATCCTATTTCATATTTTCAGAGTTGCATTTCAAACCAAAATT ATAAAATTGAATGATTTTTTTAAATATTAATTTTCATGGTGACAATTGCAGGCAGCTGCCAAGT GTAGAGCTTTGGGAGCTCGAAGTGGTCTTCCACGTCCGACAGATCCACATGTG[T/C]TATCG TTTTTACACTAGACAGAAACAAGTGGTATTTCCCATCAAAGAAGCTCAAAGCTCTTTCATCT CTCTGCTTTTGGAGTTATGTTTACTATGTGGCTCTTGTACATGTCTTGTGTAAGATGTCTT ATAGTTTGTACAGAATCATGAGTAGGGGTGCTCATGATTTGGGTATTTA
>Gm03_4 0459321	ATTTTCAATTTGTTATATGATTTGTTGTACTTGTTTTGTTTATTTACTTATATTCCATTTTCAT GGACTTTGCAGATTGTTTCTGACTCAATGAGAGCTACAAATAAGAAAATAAATGGAACACCC ATCAAGATGCTCATAGACCAAGAAATGTCCAAAGAAGTTGTTTCCAAGCACA[C/G]CCCACC ACCAAATGTAGTTGCAAAATTAATGGGGCTTGAAGCCCTCCCACAGGGTGAGCTATCTGTG GAGAGAAGCCATAGAGGAGATTATTCTCAACATATGTGTGGTCATTCAGGGACACCATTCA ACCACTGGAACCTGGAAGATAGTTTTATGGACAAGGAAATGCTTCATGAAGT
>Gm03_4 0462434	TGGATCCATCATTGAGGGACAAATACGCCAATCTCAATGACAAGGAGCCTCAGCAGCAACT CCACGAGGCCAAGCGAAGGCAGAGGAGGTCCAATCAGAAGCTTGATTTCGATTGTGTGAA TGTCTCACTAATAGAAATTAAGTTATGGATCAGAGAAGAATTACTTGATGGGTAGT[T/C]G GTTGTGCAGTGGGAGCCACAGCAGGGTCCAAGTCCCAGAAGCTGCATCTCCCCATTGGT GGACCTCATTGTGGCACAGATGAAGGAGTTAATATCTAGTGCTATGAGTTCTGTTTGGGTG GTGGATTGTGGGGACAGTAACAGCCTGGTGGTAGAGAGTGTGTCAGAAAAGAGGTTG
>Gm03_4 0537578	TATTTTTATTATTTTTATTGACGGTGAGTGATGAAAATTGAAACAGTGCAGACGACGATATC ATGGAGCTGCTATGGCACAACGGCCAGGTCTGTTGGTGACAGATCAGAACCAACGTTCTCTG AGAAAACCTCCGCCGGTGACCAACTCCCACGACGCCTCTCCCGCCGGTCTTCAAT[T/G]A CGAGAGAGATTGACCGCTGGTGGAGAATTTCAATCAGCATCTCTTCATGCACGAAGGCGA AATGGCTTCTGTTGCTTCACTATCCTATCGACGACGACGAGCCTGCGTTGGATCAAAGCTTC GGCGCCGATTTCTCTATTCTCCGCCGACGACGGCTAACAAACAGCAGCTTCATGCAA
>Gm03_4 0572180	GAACGAAAAATAAAGACTAGTGTAATCCAAAGAAACGTGACGGAGCATATATGCATGCATG CATGCATGTTTCATCATTTTCAAGGGAAGGAAGAACACCGCTGCCACCACCAACGCCAGTTT CAACGCCACCACCAATACCGCCACCGACACCGCTACCGGTACCGCCAAACCCTCCGA[A/C] CCCACCGGTAGGACCGCCGAGGCCACCACCAAGTCCACCACCTCCGCCAAAACCACCACC TATTCGGCACCAGCTCCTCCAAAGGCAGTCCATTGTTCCCAATCCCGGAATAACCACCA ACTCCACCATAGCTGATGAAGTTCTTCTGGTCTTAAAGACCAGCATCCTTGGGGTTGGG
>Gm03_4 0585266	TAAAGTGATAACAATTGAGTATACCATCTAATAATGAAAATAACGCAAATAACATAATGCTT AAAGCAACGTTTCATATCAGAAGGGAAAAATAGTTTGAATCCCATCCAACAAAAATCATGCGA TAAATCCAATTAATAAAGCATAAAGTAAATAAGTTTTTAAACAGGTATCT[C/T]GAGGAA AGCAACTACATTTAGTCCACTTCTCAATCTTGGGGCCAGCACCCTCCACCAGAAGGAG GAGCAGCATACTCATCTCTGCTGCACCAGCACCACCCACGTCTGGACCAGCACCACCTT GGTACATCTTGGCAATGATTGGATTGCAGATGCTTTCCAATTCCTTCATTTG
>Gm03_4 0585499	TCCACCAGAAGGAGGAGCAGCATACTCATCCTCTGCTGCACCAGCACCACCCACGTCTGG ACCAGCACCACCTTGGTACATCTTGGCAATGATTGGATTGCAGATGCTTTCCAATTCCTTCA TTTTGTCTCAAACCTCATCTGCTTCTGCAAGCTGGTTGCTGTCTAACCACTGGATTG[A/G]TT GCTCAATTGCATCCTCAATCTTCTTCTGTCAGCCGGGTCAAGTTTCTCACCATCTTGTCA TCCTTCACGGTGTTCCTCATGTTGTATGCATAGTTTTCAAAGCGTTTTTGGCCTCAACCTT CTTCTTGTGCTCTTCATCCTCAGACTTGACTTCTCAGCCTCTTGAACCATCTT
>Gm03_4 0587775	AATAAGTATAAAAAAATACATAAACAGAACAAATCGAATGGAAAAGATAAAATAATTTTGTTA AGAGAGGTTACCGAAGACGGTGTTGATGGGGTTCATGGCGACTTGATTCTTGGCCGCATC ACCGATGAGACGCTCGGTGTCAAGTAATCCGACGTAAGACGGCGTGGTTCTGTTCC[T/C]T TGGTCTGTTGGCGATGATTTCAACGCGTTCATGTTGCCACACACCGACGCAAGAGTAGGTG

	GTTCCGAGATCGATTCCGATAGCAGGACCCTCTCCTTTTCCGGCCATGGTTTCGGCGTGCG GCGTAGAGAAAAGAAGAGATCTAAGAAAGAGGAAGAAGAAGATGATAACCACAGCGATA
>Gm03_4 0623046	CAC TTATATGTTTTCTCCACCCAATTACATCCAATGATCCAAGGAATCTTCTTGCGACTCTT GCTCTATAAGAATACCTATTCTTCTAACACTCGAGCACACCAAACCAACCTAGCTAGTTTT CATCACCTTCCTATGATTGTTTTGTTTTAATTTCTAACTAACTAACACAGCA[G/T]GAAGCA CTGAGTCTTTGGCCATGACGTTCAACGCGAGCACCATCACAAACGGCGTCGGAAGGAGCCT GGCAGGGCGATAATCCCCTGAACCACGCTCTTCTTTGTTGATCGTTCAAACCATCCTCGT AGTCTTCGTGAGCCGCACACTCGCCTTTCTCCTCAAACCCCTTTCGTCAACCTA
>Gm03_4 0636181	ACTCCCAAGTCCCCCACCCTCCCTTAGTATACTTGAGAGTATTGCATGATACAAACCTTAG AGAAAGCCTCCTCCTGTCCAGGCAGCACCTGACTTGTCAATTATGAGCGCCACCTTTATTATT AGATGCACTCCCATAGATATCATCAATCCATCATCTTTGGCATTATCCCATCCA[G/A]TCCA ACCATCACTGCTATGGGCAGCTGACCCTTTAGCTGGTCTTTCTTCTTGTATCTTCCTGGT CCCAATCATCCCAAGAGCTTGAGTTATAATTATTATCAAACCTCCACCTGATGACTGAGGA GCTGATCCTTTAGCTGTTTTCTTCTCTAGAACATTGTTGGCCCCAATCAT
>Gm03_4 0661741	ATTA AAAAATCAATGAAGCATTAAAGAGTCAACAGAACCGAGACCGTAACCTCTAATGAGAGA GAGAGAGAGAGAGAGACTGACAGTGTTGAGAGTGCTCTGCAAATTCTGAGCGGCACCGAA GGCGAGAAAAGACCAAGAGAAAAGGCCAAGCTCAGGATGTGAATATCCGTGGAGTGGTG[G/T] CTCTTGCGAGCTTCGATGGCCGAATTATCATCGGGCACAAGTGGCGTTTTCTTCGTCCCCAG CAGGTAGTGAGACGCCATTGCCATTGATTGAAACTCTTGAGTTCTCGATTGCAAGATGGT TGGCTATTAATATTTCTAAATTTGCTTCGTCAAACGTCATCAGAGCGTCAACGTCC
>Gm03_4 0670148	CCCCTGGAGAAAGTGTGGCGTACAGCGCTAGGGTTTTGAAGCAAAAGATAACAAGGATATA GACAGCGATAGAGATAGAAAGGAATGAGAAAAAGAAGAGAGAAGCGAACCTTAGAAAGGA AGAGAAGCGTTGGGAATGGCCGCACTTGTTTCGAGTGGCGCTGTGGTTACCTAGAAAG[A/T] G]GATGGGAGCACAAGCGAGACTAGAGTTTGTTCGAGTGCCTATTCTAAAATCTAAATCTG TGGACCTATAATATACTGGGTTGGCCGATCTCAGTCCACTCTTGGGGTATTGCTATTAGC AATTCAAAGTTTGCTATTGGTAGTGGGAATAGTAGTCCCTTACTCTTATTTTGTGCGG
>Gm03_4 0676583	CCTTTCTAAAATTTAGTATTTTCTCATTGAGCAAAAGCATACATTGCTTACAACTATACCT TCAAACCTTTACTAACAGCCAAGTAATATAAATCAGATAACTTACAGGCAGAGATACAGAGCG TGAGTAAGTTTGATGAGCAGAATGCTCATAGAGTGGTTCTGTGACTGAAGTATC[T/G]GCCT CCTTATCAGACGAGCTTGAAGGCAACATGGTATTAGGAATTGAAAGGTTGCAGATTTTTTC AAGCAGCCCATATTTACTTTTTGTGCGTTGATCTTTTATTGCTTTACATATTTTTTCTCATTCA CATTATAGATGTCACTCCCCTCAGAAGCAAGTAAAGATTTCTGAGCATCT
>Gm03_4 0676856	TGATCTTTCATTGCTTTACATATTTTTCTCATTACATTATAGATGTCACTCCCCTCAGAA GCAAGTAAAGATTTCTGAGCATCTTGAATGTGTAATGACTCACTAAAAGTCCTTCCAAGATT ACTTATGAAGTTGTCGTATGCACAACATTGGTATTAGAGCACGTGTCCATCAA[A/T]TGAT CAGGTACTAAGAGGCAACTAAAAAACAACAATCAGATCAATGAGAATAATAACACAATTA TCACTAATCTACATAATAAGCATTAAAAATTAACACATTCCACAGTTATGTTTCATTATTAATA CATTGTGGACCAAAATTTAAAAAAGTTTTAAATGATGACAGATG
>Gm03_4 0678154	CTTCATACATTACTAACATAAGTCGGAAGCAAATGATACCTAAGCTTGAAAGTTGAACAAG ATTACCAAAAAATATACCTTATAAACGACGTTTGGAGCTGCAACTACAACACTATCAAAA CAGGAGCAGAGAATAGCAACCAACACAGAAAAGAACAACAATAGCAAACCTCGA[T/G]AC CGATCCGCTGACGAATTAATCACAGCTCGCTCAGAGCAGAACCAACAACAACGACGACCAA AAGAGAGCGAGAAAAGAAAGAGAAGGTGGAAGAAAACGAAAGACCTAGTGTTAACTAAGAT TACCACGCCGATCACTTTTTTCAAGATTTTTCAGACGATGAAAAGAGGATCCACCA
>Gm03_4 0678493	CGATGAAAAGAGGATCCACCATGGAACCGGTCAAAGAAGGAAGTACGAGGAGATCAACG GGCACC GAAGGCATTATTCTATGCTCTTCGAAACCCCTACGACTCACACGCCCAACAAA TGAAACCTAAACACAACCGCGCTTGATTGGAGATCGCACGCCCTAACATGAAAATGAT[G/A] GGATCAGATTGAGATTGAGATTTCGAGATTTCGATTTCACCTTAAACCATAGAAACGATCTATC TGTCTATCTATCAAATCAATTTCCCTTTTGTAGTAACAAAAATCGACACAACGGAGAATGGAG AGGAATAGGGGATCGGAGAAAAATGTGTGCGGCGAATAACGAACTGGATTAGGT
>Gm03_4 0678603	GCCCAACAAAATGAAACCTAAACACAACCGCGCTTGATTGGAGATCGCACGCCCTAACATG AAAATGATTGGATCAGATTGAGATTGAGATTGCGATTTCGATTTCACCTTAAACCATAGAAA CGATCTATCTGTCTATCTATCAAATCAATTTCCCTTTTGTAGTAACAAAAATCGACA[C/T]AACG GAGAATGGAGAGGAATAGGGGATCGGAGAAAAAATGTGTGCGGCGAATAACGAACTGGA TTAGGTTGATGTTGTCCAGATAGAGGAGAAAAAATGGACTTATATATAGGAAGAGAATCTAG AAGGTCAAGGAACCTGTTGTATTGTCTCTCTGTTAACGAGACCGAATCTCCAAC
>Gm03_4	TCGGATCACAAATCAAAAACAAGGCATATTA AAAAACGGCATTATATTATTTAAAAATAAAGC

0682560	TTTAATTTTACATCTACACCCTCAAAAGCAACAAAAAGGAAAAATAAAACCTAATTTAAAAAT GCGAATCAAATCGGGGTCAACAGAAAAACCTAATCGAAACACAGGAAACAAAA[T/A]GAATC AAGAGCTGGTGAACCTTGGTGACGGCCTTGGTACCTTCAGAGACAGCGTGTTTGGCCAACT CCCCAGGGAGGACCAAACGCACCGCCGTTTGGATCTCCCTGGAAGTAATGGTAGGCTTCT TGTTATACCTAGCCAATCGCGACGACTCCTGGGCGAGCTTCTCGAAGATGTCGTT
>Gm03_4 0683015	CCACGCTCTTCTTTGTTCTCTTCTTCTTCTCTCGCTTCCTCCTTCCTTCGATATCTTCTTCT CGGCCCTTGGGCTTCTTCTCCGCCGGAGCCTTCTCTGCGGGCTTTTTCTCCGCGGGCTTCTT CTCGGCCCTTTGGGGGAGCCATTGGTTGGTGAAGAGGGGAAAGGAGAGGGTTCGGTTT[T/G]TT ATGATGGGAGAAAAAACTGAATTAGATAGAACGAGTGAAGGAACAGAAGAATGTGTGTGT ATATATAGGAAGAGAGAAAAGCTTTTGATTGGAAGGGGTTGAGTCACGCGGATCGGTGACGT GGCGTCTTGGATTGACATTGCGTTTTTTCAGCGCCCAAGGATTGCTTCCGTCTTTT
>Gm03_4 0685818	AAATCTAGGTCTCGCATCCTTTACATGTAATTTAAAGCTCAAACAAATTCCTCGATGCGGGC TTCTGAGATTTAGACTCTGAATAGCAGGGGAAATCAATCTCTGCTGAACACGGGAAATGAG ATGTTGGCTATAACCTCTGCTCGCAACACTTTGAACCTGAATTTCCGATGAAGAAT[G/C]ATG TCTTATCTAGTACCTGTGAGGCTGTGACAGCAGGGGAAAAACATGAGACAAGTCAAACGGGT AATAAAAAATAAAAAATGTGAAAAAGTTTCTCAGTTGCATTGAGAAAACTGTAAAAAAG ATCTCGCAATAAAGAACAGCATAACACTAAGACTAATCACATAACGGAATGTCA
>Gm03_4 0689982	TCAGGAAAAAACAACAAAAAGTTCTCCAAGCTGGAATGATAAAATTAGCCTATTTTAATTTG AATAAAAAGACATAAAACAAGAAAATAATTTCAACCTATTTAAAAAAGCAACATAAAATAATA TAAAAAACCATTAACACTCTTTCTAATAGAAAGGGTGAGAATATAAAATTATAC[T/G]AGATTAA AACATTTGATATTGGATTGTGCATGTGCTATAAGAGGCAGGTAAATTTGATATTGCAGGGAT GACATATTTGAAATTAAGCACAGTTTGTAGAACATGCATGAGCTAAGAGGCAGCTAAAGTAA AATGTTTTTTTTTATTCTACATAATATTGATAATGCAGCCTTTTTTCA
>Gm03_4 0690032	CTATTTTAATTTGAATAAAAAAGACATAAAAAACAAGAAAATAATTTCAACCTATTTAAAAAAGCA ACATAAAATAATATAAAAAACCATTAACACTCTTTCTAATAGAAAGGGTGAGAATATAAAATTA TACAAGATTAAAACATTTGATATTGGATTGTGCATGTGCTATAAGAGGCAGGT[A/T]JAATTTG ATATTGCAGGGATGACATATTTGAAATTAAGCACAGTTTGTAGAACATGCATGAGCTAAGAG GCAGCTAAAGTAAATGTTTTTTTTTATTCTACATAATATTGATAATGCAGCCTTTTTTCAAG CAAACATCTTCCATTGGTCCGTCAGTTAAAAATCCTTGACGTGCATCA
>Gm03_4 0760791	TCTATTGTATGTCAAAATAAAGACCTGGCTTCCTTCTCTACTGAAAGTTACACAGTGTGGAT TTCCCATGCTGACACAGGTCACATGCCAGATAACTCCATCTACAACCTAGTTCTGATTTAACA ACAGCATTATCCTTATTTGCATGTAATTTAGTAGGCACATCTGAGGCTTTAAGAA[A/C]TGGT TCCCCCATATCAACTCTGACCTGAAACAGATAGCATTATTGTCTATCAAGCTGAAGAAAGAG AACCAATACCAGATAAGAAACAAAAACAAAATCCCTTCCTGACGTTAAACCAATTAAGGCAC CTTAAAGCAACATGGGTGATGTCTTTAAACAATATCACATATTAATAAGT
>Gm03_4 0761867	ACAATTCGTAAGAAGAACTACCTCAGGCTCACTACCATCAGAGTTAAAAATCCTCATGGTAT AATCGGTGCCACTGATGCCAGGCAAGACAAAGATAACTCCGTGACGCTCCAACGCCGAAGTT CCGATCACACAGTTGCACCGCTTTCTCAGCACTGATCTTGGGCTCGGAGGAGTCTC[A/G]G TTGTCAATCTATTAAAAAATCACGCGTCAAAAGCGAGCCATTATTTATTCGAAAAGGGAAAA GCGAAGCGGAACGTGGACTACCAAAACGAAGTCGTTTCCGAGGCCGTGGTACTTGGCGA AATGGAGGAAGCCGGACTCCTTGCGGTTGAGGAAGGAGGCGGATGAAGTTTTGACGG
>Gm03_4 0771977	TGTAACAATGAACCTGTTCTTTTGTGATGTTCTATCATCCAGTTTTGTTCCATCAACATGAC AGAACCTGAAATGATATTTAATGCAATCCCTTACCTTTATAATTTCTCTAATATTTCCCTGCA GAACCTCATAATGCAGCATGTTGATCATTCTGCTGGCGATTTGATTGACCTCC[A/C]GCAAG GGCTCCTAAGGTACGACCCCTCAGAACGGCTTAAAGCTAAGGAAGCATTGAGACATCCCTT CTTCTTCACAAGAGATACTAAAAGATATGGCTACCTTTATAAGTACCCCAAGCATAATGGC TGGGAGGAGAGAAAACACAGTTGAGAATGAATTGCATAGAATCTCATACTTG
>Gm03_4 0785291	GGCCTTACCCATGGATTCTCTTCCCAAATTGCGCCACCCTTCTCCATTTCCCACTCCCTTAT CCTCTTCCCTCTCCTTCCGCTCTTTACCGCCACCCCATTTTCTTCTGTCGTCGTTTAGGTTA CCCTCCATCAGAGCCTCCTCCTCACCCCTCCAAAACGACCCCGCTTTTCGCACCC[A/C]AAG AAACAAGCCCTCCCTCCCCCCCCCTCTCAAACCCTAACCTCCTTCTCTCCCCCTCCTC GAAACCACCTGCATCGTCATCGCCGCCGCCGCTTCTTCTTCATGCGCTTCCACCACACGC CCGCCGTCATCGCCGCTCCGCTCACTTCCCCCGCCGCGGAAACCGACACCGCCTT
>Gm03_4 0785299	CCATGGATTCTCTTCCCAAATTGCGCCACCCTTCTCCATTTCCCACTCCCTTATCCTCTTCC CTCTCCTTCCGCTCTTTACCGCCACCCCATTTTCTTCTGTCGTCGTTTAGGTTACCCCTCCAT CAGAGCCTCCTCCTCACCCCTCCAAAACGACCCCGCTTTTCGCACCCCAAGAAAC[T/C]AG CCCTCCCTCCCCCCCCCTCTCAAACCCTAACCTCCTTCTCTCCCCCTCCTCGAAACCA

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>Gm03_4 0824534	GGAAAATCCAGCCGCGGCGAGGCGAGGGCGGTGTTGAGAACTTCAACCGCAAATTCGA GGAAGCGTATCGGAAACAGAACTCATTGTCGTAGCTGACAGAGAATTAAGAGACGAAATA AAAGGTTGATCGCGAGAAGTATAGTGCCGAGATATCGCGAGTGGTATAATGTTCTTC[A/T] AGCTAAGGTTGGATCAGTGAGGGACTTGACGGCGACGAGTACGTGACGTTTACCCCTGA GGATATTGAAAATTACCTCGTTAACCTCTTCTTTTTTGGAACTCATCCAGTGCTGTATCGTC CTCGTCGGTATCGTCGTCCCCTCTCCGGCGTTGGAGTTAAGTGCACAGTTTGAACGG
>Gm03_4 0824686	GATATCGCGAGTGGTATAATGTTCTTCTAGCTAAGGTTGGATCAGTGAGGGACTTGACGGC GACGGAGTACGTGACGTTTACCCCTGAGGATATTGAAAATTACCTCGTTAACCTCTTCTTTT TTGGAACCTCATCCAGTGCTGTATCGTCCTCGTCGGTATCGTCGTCCCCTCTCCGG[G/A]GT TGGAGTTAAGTGCACAGTTTGAACGGCGAGTTTTTTTCGCCCTCATTTTTCTTGATTGTAGT AATGAGTATAGAGTTTTGACATAGACAACAAAGCTATACCGACTTTTTTCAGTTTTCTGCGT GTGGTACACGTAAGCATGACTTCATCGTGTAGTTTCCACTTTTTACTTTCAGAC
>Gm03_4 0843491	TCTTCTGCAAGAAATGGCTACATGGAGTTGTCAAGGCCCTTGTGAGCAAAGAACCAGAAA TTGCAATGAGAATTGATAAAAAGGGGCGAGACAGCACTCCATATGGCAGTTAAAGGACAGAA TCTTGAGTTGGTGGATGAGCTCGTGAAATTGAATCCATCTTTGGCCAATATGGTGA[T/C]A CCAAGGGGAACACTGCACTGCATATAGCAACCCGGAAGGGTCTGCTACAGGTAACAAAAAC TGCTTTCTTTTCTGTTTTCTTTCTTAAGAAAGTAAGTGAATGTATTGTTCTATGTCAACTG AACTTGTGCTTTTTATTTAGACATTAGATTCTAGAGTTCCTCATCATCCAAGAC
>Gm03_4 0844367	TAAACACAGATGTTATCAACAAATCTGGAGAACTGCTTTAGATACTGCAGAGAAAAATGGT CGTTTGGAAATTGCCAACTTTCTGCAACATCATGGAGCTCAAAGTGCCAAGTCCATCAAGT CACCTACTACAAACACGGCCCTTGAGCTCAAACAAACAGTGAGTGACATAAAAAAGT[G/C]GG GTTTATAACCACTGGAACACACAATTAACGCAAAGACGTATGCAAGGTATAGCGAAGC GAATCAACAAAATGCACACTGAGGGGCTTAACAATGCGATCAACTCCAACATTGTTGTTGCT GTCTTATTGCAACAGTTGCTTTTGTGCTGCCATATTTAATGTCCCGGGCCAGTATC
>Gm03_4 0870653	GAACCCCTCGTTCTGCGTCCTGGACTCCCACTACATCCTCCTCCTTTCACCTTCCGCGAC CTCTCCCCTCCCCCGTTGCCTCCTTCAACCGCTCCTACCCCGTCACCTCCCCCAACGAAT ACTCCCTCTTCTTCGCCAACTGCGCCCCCGAAACCTCCGTCACCATGTCCGTCCACA[T/A]C GAAGCCTACAATCTCGACGCTGACGGTTCACGCGACTACCTCTCCGCCGGCCAAACCCCTC CTCCCTTCCCTCTACTTTCTCTTCTTTTAACCTACCTCTCCTTCTCTCTCTGCTCTAC ATCTGCTACACCAACAACTCTCCGTTCAACCGCATTACCTCTTAATGTCTCTCCT
>Gm03_4 0874471	ACATAAGTGAAGAAAAAGAAACCAAAGTGATAGATAGCTATATCACTAATAAATCTTCTCTC TGGATCATAAATAGGAAGAAAAATATATGTAAAATAAAACAAAAAGTTCAAACCTTTGATTAA AAATCAATATTTTTCGTCGACAACATCTACACTAGCGGAACAGGACAAGACCC[T/G]TCGGA GAATCCCAGACAAAGCAGAACAGAGACCTCCTTAGCCTTACTCCAAACCAAGCAGCACTT GGTTTTCTTAGCCTTATCATTAGGCTTTAAAGTCTCGTTCCCAATATCTTTCTTCTTCAT CTTTGCCCTACCAAAACATTTGTGGGTTTTCTCGTGCTCGTTCTCTTCTTC
>Gm03_4 0885382	ACTGGGCAAGCACATTCTACGATGCATGAGGCGTTCCAACAGTTGAAAACCTTTCCCTTTC ACACTTTTCTTTTTCTCTCACTTTCAGTTCTCACGAGAAACCTACGCCCTTCCGCTCATC CTAGAGCTCCCACACCCAACAACCCCGATTTTTCTCAACACACACTCACACAATC[T/G]GT TTTTCACTCCCTCGATTTTCGCGATCTCGCAGTCGATTCTCCCTTTCTTTTAATTATAAACGC CGTCGTTTGCTTCCCCGCGAAGTAGTTAGGGTTTCTTGTTGGCACAATTATTGGATCGCTG CATCGATAATTGACGGGAGAAACAATTTTTTTTTAATCGAGAAATTGTTTTCTC
>Gm03_4 0886333	TGTGGAATTAATCAGTTCCACTTTGCGATTGGAGCAGGGTTCTTTGACTGGAGAGAGTGAA GCTGTGAATAAGACTAACAAGCGTGTGGATGAGGATGCTGATATTCAGGGGAAGAGGTGTA TGGTTTTTCGAGGGACCACTGTTGTTAATGGGAATAGCATATGCTTGGTCACTCAGA[C/A]T GGCATGGATACCGAAATCGGGAAGGTGCATATGCAGATACACGTAGCCTCGCAGAGCGAG GAGGATACCCCATTTGAAGAAGAACTCAATGAGTTTGGAGAGAAGCTGACCTTGATAATTG GACTGATATGTATTTGGTTTGGCTCATCAATGTTAAGTACTTCTTTCTTGGGAGTA
>Gm03_4 0889026	CTCCTAGGGAGGAAGTATACCAAGCAATTGAAGACTGCAGAGATGCTGGAATTCGTGTTAT GGTTATTACTGGAGACAACAAGAACACCGCTGAAGCTATATGCCGTGAAATAGGTGTATTTT CTCCAGATGAAGACATTAGTTCTAAAAGTTTAACAGGTAGAGATTTTATGGAAGT[T/A]ATG ATAAAAAGGCATATCTGAGACAGCATGGCGGGCTTTTGTTCGAAAGGGCTGAACCTAGGCA CAAGCAAGAAATTGTGAGGCTGCTCAAAGAGGAAGGGGAAGTGGTGGCCATGACTGGGGA TGGTGTTAATGATGCTCCTGCCTGAAGCTTGCAGACATTGGTATTGCAATGGGCA
>Gm03_4	TTAACAGTCACTAATCTTATTGGTTCTGATGCAACGACTCTGCAATTTCAATTGGCTGTAGTT

0906651	TCAGTAGTCAGAATTGTTTTATAACATCTTTGCTTGTAAGCTGAGTTTTGGTTACCTTGGTTC TTGCCAGCCGATCGAGATCTCCTGTCAGAAGAGAACGAAGCCCAGTTTCTGAA[A/C]ATAGA TCACAGAGTCGTGAGCCCTCCAAGATAAGGAAGCATAGTGCATCACCTGACCAAAGCAGC CCACGGAAGAGAGGTGACACATCTCCTGGCAATGATAGGTTGGCCACTCATCAAGATGGG TCTGACTACAGTGATGGTCCTAGAGGGAAAAGTAGAAGCCCTGCTAGCCCTGCAA
>Gm03_4 0915539	ATCACCGATTCAAGTTAGGTGGAATCAACTTGAGAAGACTTTGGTTTCCAATTGGTAGGCCT TGGTGAAGAATGCTGGCCAACAGAGGTATGTAGCCATTATGCTGCTAACAGTTGCTGCACC AATCATCATATTATCACCACAATCTGTAGTTGAATTGCTTCCAACGGCGATGCCC[C/T]ACC CATAATGAGACCAGTCATTGCCCCAGGAAGGGATATTAGACCCACTGTTTTGGTGTGTGCC ACCACCGGAGAAAAGGGCCAATATCAGAGCCCTTTTCACTTGTTTCATGCGTTGCTTGCTGTG GGGTTGCACCAAGAGACAATGCTGTCTCAACCTGAAATACAAATTAATACTCT
>Gm03_4 0915692	TTGAATTGCTTCCAACGGCGATGCCCCACCCATAATGAGACCAGTCATTGCCCCAGGAAGG GATATTAGACCCACTGTTTTGGTGTGTGCCACCACCGGAGAAAAGGGCCAATATCAGAGCCC TTTTCACTTGTTTCATGCGTTGCTTGCTGCTGGGGTTGCACCAAGAGACAATGCTGTCT[T/A]AA CCTGAAATACAAATTAATACACTCTTTCATTTCATCCCTCTCAATTTACTCATCCACACCTATT ACAAATAAATGTTAACCCCTAAAGCCTATCAAATCACTTGCTGTGTGTTTTCAACCTTATTAT GCATGAAAAAAATATTTCACTAAAAATGATTTGTGAGATTCTAGTCATTAA
>Gm03_4 0942035	ATTACTAGGGCATCCAAATCAACACACTAATAGATAGATTTGGAATTGGACCTTTTCGGCTCG GGAATTGCGCTTCTCTTCTCGGACAACCTGGACGGAAATGCCGAATCGCTCGGCGCGGCG CATTTTCTTCTGAATATCGGAGAGCGGAGCGTCGTTTCCAGAATCGGGCGGCGGGGA[A/T] GTTTTGGCGTCGGAGTCCTTAGTGTCGGCGGCGGCGTCGTCCTCGGTGGCAGGGATTGGA TCTGGTTCGGTGGGGTCTAGGGTTTTGTTAGGGTTTTCTGTGCAGGGTTGTTATTGTTGT CGGTAGGGGTTGTCATTGGAAGCGGGAGATGGATCGATCGAACGATTAACGATTTCGGAA
>Gm03_4 0951123	ACAAACACGTTTTGAGACGCCGTTTCCACGACCTTGCGATTGAGCGCAGCATTGAAGGCGA TTTTTTACTAGCATGGCAGCAATAATGTTGACGACTATGAATATTCTCTGCACACTTCGGT TTGGGTTAATTCCCGGCGGAAGAGTTGGTGCTGCTACGCCTCACTTTCGACGCCAG[G/A]T AGGAAGGAGATCCGTCTCGGTTTGCCAGCAAAGGTGCGATGTCCGTGACACCCCTCGAA CTTCTCAAGGTCATTCCAATTTGCACTCGATTGCGCACTGTTATCTTCTACATTCTACACTAC TAATAACTCTGCATTCTTCTTCTGTGTGAAGAATTGTCAATTGTCAGTGAAGCA
>Gm03_4 1007691	CAAAAAAAGAAGCCCACTTAGAAGGACTCTTTAAATATATTTATATTCAACGTATAAAACCTT TGATGCTGTTCTGCCAAACCCCAAACCTAGTTTTGTCAGTAGCACACTTCACCCATGGCA GTAGCAGCAGCTCGCCCTATCGTCTCCGTCCAAACCCCTAGAGGGCGATGCAACCC[T/G]TA CCGTTCTCTTCCCGACGTCATGAAAGCCTCCATTGCGCCTGACATCGTCAATTTTCGTCCA CTCCAACATTTCTCGAACAGCCGCCAACCCCTACGCCGTGAGCAAGCGCGCCGGCCACCA AACCTCCGCCGAGTCCTGGGGCACCGGCCGCGCTCTCGCGTATCCCGCGAGTCCC
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>Gm03_4 1008255	GGAATCCCCCTCGTCTGTCAGCGACACTGTGAAAGCGTCGAGAAATCCAAAGAAGCCGT CAAAATTTCTCCAGAAAAATCGGCGCATTACAGACGCGGAGAAAGCTAAACTCAGCCACGGA ATTCGTCCCGGGAAGGGAAAAATGAGGAACCGCAGGTATATCTCTGGCAAGGGTCTCT[G/ C]TATAGTGTACGGAACAGAAGGTGCTAAGGCCATTAAAGCCTTCAGGAACATTCCCGGTGT AGAGGTGGCAAACGTTGATAGGCTCAACCTTCTGAAGCTGGCACCCGGTGGCCACCTTGG AAGGTTCAATTATTTGGACCAAGTCCGCATTTGAGAAATTGGACTCTATCTATGGATCCTT
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>Gm03_4 1008459	TAAGGCCATTAAAGCCTTCAGGAACATTCCCGGTGTAGAGGTGGCAAACGTTGATAGGCTC AACCTTCTGAAGCTGGCACCCGGTGGCCACCTTGAAGGTTCAATTATTTGGACCAAGTCCG CATTTGAGAAATTGGACTCTATCTATGGATCCTTCGATAAGGCCTCCGAGAAGAAAA[C/T]T GGCTACTTGCTTCCCAGGCCCAAGATGATCAACTCTGACTTGGCCCGTATCATTAACTCCG ACGAGGTCCAGTCCGTGGTCAGGCCCATTAAGAAAGATGTTAAGAGAGCACCTCTCAAGAA GAACCCTCTCACGAACCTCAATGCGATGTTGAAGCTCAATCCCTATGCTAAGACCGC
>Gm03_4 1063028	CATAAGATTTGATGATTCAGATTACATAGTTATCTATGATACACATGTTTTTCCGCTGTGGT TTGCAATTTTATATCGTAATTTTAGGAATTGATGTTATCACAAGTAAAAGAACTCTTACAGA TGATGAATTTGTGATGTTAATTGTTTTTCTTCTTTGAATCTTCTCAGGGT[A/C]AACATAT GATGGTTCCAATGCAAGGCCCGGAGTAGGTATCAAGTTTATTATTGACCCTCTTGTTGTAAC GTATGTTTTCTACGCCAGTTACCAAGTGCTCACGGCATATTGAATGCTTTTTATGTTATGTG AATACTGACAGGAGATGTTGGTTCTTGTGTCCGTTTTTTTTTAAAT
>Gm03_4 1082727	ATGATGTAGAGACAAGAAAACAGACTTGATTCTAGAAACAAATACAGTGGTCTACATTAGT TCTCTGCGCCATTAAGAAGCCGCTCCATTACTGTAGGTTCAACATATCTAACTACAACACAA TATCCGTCACGAGTTTATCACAATTTCCCTGAGGAACTGCTTCAATGTTTCATGAC[T/A]CATG AACTTGATAACACAAAATACCTTGACCGAAAAATCGTAACAATGAAACCAACCATATAAAA TTAAATCCATTTTAGCCAAGATATCAAAGTGAGGCAATGATTACCAATACCTGTGCCGCAT CTAATGATGGTTCCTCCTCCACAAAACCTCCAATCTTCTTCATCATGTTTGTA
>Gm03_4 1095513	GTGAAAAAAGGAAGGGGGGGATAATTAATTAATTTATATTGCTTTGAATCAAAATTGCAACC CGTACCAAGTATCCACGAAAAGCTGTTTGAATTTTGATAGCTGCCACTTCATCCTTTGGTTT ACCAGCATAATGAGCAATTGGGGCAGCTTCAACCCTAACAGGTTCTATCTGAACA[G/C]AAG GAACAGGTTCCCTCGGCATCCATGGCAGTTGCAACTTCAACAACATAGCTGTCATCATGACT GATTTCAATTTCAATATCAGTTAATTTAATCGCTTCTGGTGGAGGAATAGGTCGCGGTGGTG GTGGTGCTTTATGAGCTTCTGAGTATGAGACTGAAGTCTCCAATTTTGCTTCC
>Gm03_4 1170229	TCAGAGATTCTGTGTTACATAGCCAATCCTCTCACCAAGTGCAAGTAGTTCCTGGTCAAA CCAGTCATCAAAATAATTAATAAATCTGAGTGCATAATTAATTGCAACGCTAGAAACAGTA ATAAATTAATAATCACCTCATAACTCATGTTATCTACGTCCATCCTCATGTCTC[T/A]GTGCTG ATCTAGCATGTTCTGGAACCATACACTGATGCACGCTCAACAACCATGAAACCCTGAAAG CCAACAAAAAATACAAAGAAAAAAGTAAATGAGTCTCAGCCAGTGGTAGTAATGCCAGAA AGAACAATTTTGATGTTCTTAATGAACAGCGCAACTAGTTCAAAAGAATATA
>Gm03_4 1171468	TTCCGCAAAGAACCCTCACCTTTATTGAAAGGCCAGTACCTCTCAATGTGGGTGTCATAGT ACAATAATGATCCCAGGGAATATATTGACAATCAATATTTGGCTTCTCCTGCCACAATTCTG AAGATATAGGAAGATCAGTTGAACTCCAGCATTAAGTATCTACTAGTACTGCC[A/G]CCC TCATAGACTGAAGGAATTCCAGGGCTCTTACGCTTGATGCCCTTGTCACCTTCCCATTGT AAGATCAATAAAACCACCATCAACATGAAATGTTTGTCTGTCATAATTGGAGGAAGAAGGTT GGTGTGCATAATTTTCCAACGGTATAAACAATGATGGTCCAGCACTAGACAAA
>Gm03_4 1184564	TTTAACAAATGGATCAAGCTGTCCACTGATCTGTAACCGTGGTGGACTCGACAAGACAACA ACAATATTCAGAACATTAACAAGACCACACCACACCATGAGCGTCATCGACATTCTCACCAG AGTTGATTCCATTTGCAAAAAGTACGACAAATACGACGTCCAAAGCCAAAGGGACT[A/C]TA ATCTCTCCTCCGACGATGCATTGCGCCAACTCTACGCCTCCGTGACGCCGACATTGAGGC CTTACTTCAGGTACACTTCCCTTCACTTTTTTTTTCAAATTCATTACTAAATGAGGAAATTT GTGCCCTAATTCGTGAGAATTATCGATTTTTTTTATTCATCTCCCAACATGTG
>Gm03_4 1207607	AAAGTTTGCTTCTTAGGCTCATCAATGGGGTTGAATCAGTTCACTGTAGGCCATCAATGAA GTCCTTTCTGACTATAATTAATTAACATATCATGACATATCTAGTGTGGTCTCAATTCATG TGTAATAACACACCAAAATTATTGGGAATTAGTTGGTGAAGGTTAATTGGGAC[A/C]AAGTT AGAGAGCGAGTTCGAGATCAGAGATGCATGGTTCTTTGTGCGTGTTGTTAGCGGTTCCGAG GGTAAAGGGACCCACTGCATGTTTGAGATGAAAGAAGAAACAGCTAGTTCCCCGAGCAGT GCATGTTCTCTAAGTAGTAGTTGTTCTTGTGGTTGTCTTCTGTCTTAACCT
>Gm03_4 1223681	TATCAGCTATTTGTGAGGCTTTGAGGAGTTTGATGAAGGTTGTGAATTGTCCTGCTTTCTCA AGCACTTGGGTGATGTTGGTGGGTCCCGCTGGTGCCGGTGCTACTTGGGCTGAAATTGTTT GGATGAAGATGTTGAGGAAGAGGAAAGGGAGGAAATTGAATAGCTTTGTCATTGGT[A/C]TT CCTTGTGAGTGAGGAAATTAAGAAGAGAAGACTAAGATGTTTGTTACAAGTGGAAGGAAGA GATGTGGTGTGCGTAATATATATTAAGAAGAGGGTGCATGGAAATTTGATTGTTTAAAGGTT AGGTGAAAGGTTGCAATTAGTATAAGGTAGCAAGAGGCCCTCATGCAATCAATTT

>Gm03_4 1228284	TAGGCCTTGGTTTGAGTTGTTTCAGCTGTGAATTGATTCTATCAGCTATTTGTGAAGCTTTGA GGAGTTTAAATGAAGGTTGTGAATTGCTCTGCTTTCTCTAGAACTTGGGTGATGTTGGTGGG GCCCGCTGGTGCTGCTGGTGGTGGGCTGAAATTGTTTGGATGAAAAGGAAAAGGA[A/T]AA AATGGATTAGCTTTGCCATTTGTGTGAGTGAGGAAGCTAAGAAGACTAAGATTTTTGTCA AGTGAAGAAGGGGATGTGGTGTGTGATTTATATTATATGGTGTAAAGGTTTGAGGGTG AGTGGAAATTAGATTGGTATAAAAAAGTTAGGTGAAAGGTTGCAATTTGTATAAGG
>Gm03_4 1230797	ATTGGCAGTTCCCATTCCCATTGCCATCGCCAGCGCATATATCAAATCGAACAGCCGCAAA CCCTTCCTCTGAACCTCTGATCCGCACTGGCACTGAAGCCATTGCTGTACCTTCCAAACCCCT AGCGTAGCCACGCGCTGAATTGCGATACGATGATCGTGCGCACATGAGGCGGCACGG[G/A] ]CCGGCACGATGATGAACACAGCCAACGGCACGATGATCTCACCGTCGTCTTCGTCCGGGAA CGGCGGCGCAGTCGCCGGGACTGAAGACCTATTTCAAACCCCCGAGGGAAGATATAAGC TTCAATACGAGAAGACTCACCTTCCGGTCTTCTTCATTACGCCCATGGCAAAACCGTTA
>Gm03_4 1231547	TGACAAAATGGAATTGTCTTAGCATATGAGTGAATTGTTGGTTTGATTGAATGAGCAGGTAA CCCTGGCACATCTCAAGGACAAGCCTGCGCCTTCTACCCCAACCGCTTCGTATCGTCAAG TTTCAGTGCCAGCAGTGGGGTACGGTCGGCGGCAGCTCGATTGCTGGGAGGAAGCA[G/C] TGGTAGCCGGGCACTTAGTTTTGTTGGAGGCAATGGTGGCAGCAAGAGTAATGGAGGAAA TACTAGGATTAGTTCCATTGGTGCTTCGAGTACAAGTAGTGCAATGGCGAATCCGAATTTTG ACGGGAAAAGGAACATACTTGATATTCAATGTCGGGGATGCAATTTTCATTAGTGATTT
>Gm03_4 1234338	GATGATTTAGTTCAAGTTTGGAGCATGGAAGATCGCAAGGTTGTGGCATGGGGCGAGGGA CATAACTCATGGGTAAGTTGCCTGACAATTTTTAGTTGTGTGCTTGTAGCAAACCTTTGCTAC CGTAGGGTTAATGTAGTGTTTGTATCAAATGCCATGATCAGGTCAGTGAGTTGCTT[A/C]T GATTCATATTGGACATCACCAATTCAAATGACAATGGAGAGACTGTTACATATCGTTTTGG TTCTGTTGGTCAGGTGCTCAGAAAAATGTTTCATTATAATTTTTACTTTTTCTTTGTGTAGCTT TTCATTCTAAGTAAACTCATTGGTATACCTTAACATGGATCTGTGGTAGCTTT
>Gm03_4 1236752	AATAAGACATGTGATTATTCATCTCTTCAACCATCTTTACAGTACAGAATCCCTTTTTGTTCCCC TATAATTGTACAAAAACAACCTTTCTTTATCTTTTACCTACAGGATTTGCTCAGAAGTTTCT TGCAAGTCCAATATTATAATATGGCCTCACAGTTACAATACAACCTGGCTTCGA[A/G]TCAGCG GTCTCAATACAAGGAATGCGATACTTGTGTATTCTCGCGAGTACCACGGAGACAAAAAAGG GTGTAATGTGCTTAAAGTATAGTCCATACATTATACTTTTATATACAGCCAATTAACACCAT CAATATAAAAGTATATTCTATGTCAACTCACACTCATGCACTGCAGTG
>Gm03_4 1255346	GCCGGCAGGGAAGGACAATTGTATCTCGGGGCTCTGACAAGCAAATAACACATTTTCCTTG GTCATTCTCCGTTGAATTTCTAATACCATATATCTCCTGCAGCTCATACCTCATTCCATTAC CCACAAGATCTGCTTGACAACCTTTCACCCGGAACCTCTCCTTTCTCCTTCTCAAAC[G/A]TCGT TTGCGTAATCTGAGAGTTTGTGTTACTAGATGTTGGAGATCTATCTGACTCATCATGATTAC CAGGGGATGCGTCTGCCTTTACTGCTAGAGGTAGACATCCTTTTCCACCCACCTTTCCAA TAACCTCAGACTCTTCAAACACTGAAAAATCAATACCAGTTCAGCTGCCTGTC
>Gm03_4 1275788	GAAAAGGAAATCGTGAGAAACGGGAGCGAAAACGAGTTGATTATAGTAAGAAAGCGCGTG AAATCCGAAAAGGGCTGCATTGGCGTGAAAATGGTAGATCTAAAAGTAAAGGAGAAAACCG AAATAAGAGATTGGAAGGTACCGTAGCAATCGGGTCTGTGTTAGATTAGAGAAAAC[T/C] AGACACAGTGTCTTAACGAAAACCTGGAGGACTGCTTCGCTGATTTGCTTTTCGCTTCGC CCAATCAAATCAACACAACCCACCTATTTCTTTAATAATAAATAAATCACACATATATAATTTA ATTTTACTCACAAACCATATTCTTTTCGCAAAGTGTGGACATAAATAAATGACA
>Gm03_4 1595526	CCACCAGCATTGGAGAATCATCAACTTCTGAGTATTATTATCATCTTCTCCCCACTGTA GATGTGATTATTGGCCCCACCAACAAAACCTTGCATCACCAACTCTGCAGGCTCCATGACA TGGCTTATCTGGTTGTTGCAGCTCCATCAGAAGAGCTGCAACTAGTGGCCGTGGC[C/G]G TGGCTGTGGCTTCACCTCCAGACACAAGATGCTGCTATCTTTGAATTGGCACTTGGGGTC CATGAAACTCTCTTGTGCTTGAACAAGGAATCAGCATTAGCAGAAGAGTTGTTACCACTCA CAAGACTTGATGATGAGTAGGAAAAGGAACCGTGAGGGTGGTAGTATGAGATGTTG
>Gm03_4 1895733	TGGAACCTCTACCATCGAAGCAATATGTGCTGGAATGCCAATGCTTACGTGGCCGTTGTTTG GGGATCAATTCTTCAATGAGAAGTTCATTGTGCAGGTAAGAATTGGAGTGAGAGTTGG GGTGGAACTCCTGTGAACCTGGGGCAATGAAGAGAAAAGTGGGGTCTTGGTGAAGAA[G/C] GAACATGTTTTGAAGGCTATACAAGTCTTAATGGATGAGGGAAATGAAAGGGAAGAGAGAA GAAAAAGGGCTAGAGAGCTTGCGGAGATGGCTAAGAAAGCTGTAGAAGGAGGATCTTCTC ACTTCAACGTAACCCAGCTAATACAAGATATCATGCAACAGTCCAACAAAGACTAGTAC
>Gm03_4 2001213	ACCGGACTTAAATATGTCATTCTTGTGCTGAAAAAACTTAAACTATATATTTGATTGCATAA TGATCATTATTTTCATGTAACACTTCTCTTCAAATAACCAAGTATAATAACAGTGTTTCCAATA TGAACATACAACAAAACCTCCATGCACGGGGATAATGGAAGATTGGAATAAC[G/C]CTGGCA



	CCAACTGGCTGTGTTGACCGCAGCCATTATTAGTCGCTGGAAGCAACCAATATCTTCCCAG ACCAAATCAGGATGCTTTTTCTTCAACAAAAACATAAATAGCACGCCTGAGAATTTGCACA AACTCCGTGTCCATGGTCTAAACTTTTTGGTAATTGATATATACTAGCCTCT
>Gm03_4 2060956	TATCCAGCAAAAGTCACGTGGTAAGTAAACAAGTCAACTCCGGTAAGTAAATCAGATCAATA TAAAAGGATGTTCAATTACGAATTGCTTGGCCCGTATGCATTGGAAACCATAGCACACCACA ATATTGCTCATAACTAGACCTAGATGTATACCACAAACCGTATAACGTCGTACA[G/A]CTGG CTATTGCTACAATATAGAACGTATACACTTCTTTTTACATTCTCCATGTGGAAAACTTTAACCC AAAAAATTGGCATCGAACATATTCCTATTATTTTTTAAGTTTGACATCCTGTAATAATAGTTA GCCTAACATTCATTTTCAATCACAAAATATAGCTAGTCACCAAACCCGG
>Gm03_4 2062804	TAAATCATATCAGTAATAGAATTCAAACCTGTCAGCAGGACAAGGGGGTGTCAATCACATTG ATGGACACCTAACAGAAAAAAGATGCAAATGAAACAAAAAGATCTAACCTCAAAAGCTCCA CTGAAGCTCCATTTAGCTGATTTGTCTTGAGTCGGGGGATCACAAACAGAGAGAA[G/T]TGG CATGGTTGGCCTATATTTTGTATTAATCTGAAAAGTGCATGCGATGCTTTTAGAAAAGCCT TACAGACACAAGTGAAATGCAGGGAAGAAAGATTGATTTGGACAACAATTGATGCATAAT ATTCATTTTCCAAGATACTAAATATGAACATGTTTCATATCAAATTATTCGTTGA
>Gm03_4 2139098	AGATGGCGGAGCACTTGGCATCGATATTCGGGACGGAGAAGGACAGGGTGAACGTGCCGT TCTACTTCAAGATCGGCGCGTGCAGGCACGGCGACCGGTGCTCCCGGCTTCACACCAAGC CGACAATTAGCCCAACATTGGTTCTGTGAACATGTACCAGAGGCCCGACATGAACATG[C/ A]GCATTATCACCAACCCCGATCAGCCACAACCCCAATCCCTTGACCCGACAAGGTGCAG GATCACTTCGACGACTTCTACGAAGATCTCTTCGAGGAGCTCAGCAAGTACGGCCCCATTC AGAGCTTGAATATTTGTGACAACCTCGCCGATCACATGGTCTCTCTCTCTCTCCACCCCTT
>Gm03_4 2179109	GGCAGAGCCACTGCAACAGGAAGCAGAGTTCGCCGCTACTCCGCGACCCCTACCCTATCAA TTTTCGTACTTCTTTTCGTAATTGCAGCGAAGAATGGCGAAGAGGAAGGAGCGGATTGAGA ACCCGGAGCCATTCGATCCGTACGGCGCCGACCCCGCGAAGACGAAGAAGCGTTGAA[C/ T]CCGCCGAAGCGGCACCAGCAGGAAGAGCAGTTTCTGCTCCCAAACCTCAGTCCAAGAT CATGAAGCAGGCGCTGATCCAGCAGGAAGGAGGAAGGAAGGAGGAAGGAGGAAACGCACGAGACA ACGCCGCCGCGAATTTATTCGAAGAGGTTCCGAACCTTCGAGGAAGACGGCGGTGACGACA TC
>Gm03_4 2184207	TTCTTGTAATGATATTTTCAAGGTACAAAAATGAGCTGCAGAAGGAAGATAAGGATAGATTAAG GAATCTACTGAAAAAGCAAAAGCATAAATTGGTGAGAACACTAAATATGAGTTGCTTTTTCTT CTGTTTCAAATTTCTGGTTGTGATTGATATTTTTTTTCCATACTGCAGGTTACAC[T/C]TGA TTAGTAGAGAGCTAGACCATAGCTGCAACCGAGGTGAAAAGGAGGAAGATCTTATGTCAAT TTATATCCTTTCCATCCTCCACCTTTCTTTACCCTACTTATCCTCATGTATTGAGATTGCAAG AAATTTAAAGTCAGATAATATAACTAATTAATAATATGAAGATTCTTTTGA
>Gm03_4 2223604	TCCTCGGAGCCATTCTGGGTGGCCAACCCAACCCACGCAGCATCCTGGTCCGACACCACG CGCCTCTTCTTCAACGGCAGTCTTGTTTTGTGACAGCCCGAAACGCGCTTTCGTGGTCAA CCGCCACAAACAACGGCGACCGCGTGGTGCTCCTCAACACCTCCAATTTACAAGTTCA[G/C ]GACAAAGGCGGCACTCCCTTGTGGCAGAGTTTTCACTTTCCCGCAAATACCCTCGTCCAG GACCAGAATTTACCTCCAACATGACTCTTTTGTCTCCAACGGCATTTATTCTATGCGGTT AGGCAACGATTTTCATGGGCCTATACGAAAACCACGACTCGTTATATTGAAAACGCACG
>Gm03_4 2224125	GTTGGAACCGGACGGGAATCTGAAGGGTTATTACTGGGACGGTTCCAGGTGGATGCTTAA CTACCAGGCGATTACTGAAGCCTGCGAGCTCCCCCGCTCCTGCGGTTTCGTACGTTTGTG CACGCCGGGTGGGTGAGGATGCTCCTGTTTAGATAACCGAACCCGGTTTGAACCGGGTG[ G/A]GTGTTTTAATGATGCTTCTTCTGGAGACGCAGACTTGTGTAGTAGTGAAGGAATTGGTG GAAAAAGTAGTTACTGGGTGCTAAGAAGAACCGGAGTGGAGGCAGCGCATAAGGAGCTAC TAAGGCACGTAACGACGTCGTCTCTGGCGGAGTGTGAGGGATTGTGCCAGAACAACGTGA G
>Gm03_4 2224392	AACCGGAGTGAGGCGAGCGCATAAGGAGCTACTAAGGCACGTAACGACGTCGTCTCTGGC GGAGTGTGAGGGATTGTGCCAGAACAACGTGAGCTGTTGGGGGGCGCTGTATAGCAACGA AACCGGGTTTTGTTATTTGTTGGAGTATACGATCCAGACCTTGCTGGGTACCGGGGATG[C/ T]ATCAAAAGTGGGTTATTTCAAGGTTAAGAAAGAGGAACGAAGAACGAACGGGTTTGGAT ACGGGTTGGGGTTGTGGTTACGGTTTTGTTGGGGTTGGGGTTATTATTGGGGTCGGG TTTTGCGTGACGAGATGAAAAAGAAGAGAGGGGTGAAGGAAGAGATTGGGGTTCCGC
>Gm03_4 2224778	CAAGTTTTAGATCCATTGAAATGAGCAATAGTAACAGTGCCAACGAATGACGCAAGGGGTG ACGCATGTGCTCTTAGTTGACAATTGGTGGGGTCCATCTAATCTAATGTTTGTCTGTTG GTGTTGAGGTCCGGTGAAAAGGTACCGGATTGTGTTGGTTTTCTACCAACCCAGGG[C/G]C GCAACTCTCGCAAGTGGCGGTTTTCAAAAACATGACTTCACTACAGTGTTATGATCATAGCAT

	GGTAAAGTCAATCAGCTTCATAGACTGAACTTAAACTAAATAAACATTTGTGCGACGTGTCT TTTTTATATTTCAATTTGACTTGAAACTGCCGCCGCCGCCGCCGCCGCCGCCGCCGCCGCC
>Gm03_4 2243466	GCTCAAGATCGGCCCAAACGAACCATCCGAGCTGGCTATCCATGAAAATGCCTACGGTCTT GCCCGATATGCCGTCATATGCCAGGAGAATGGTCTGGTACCTATTGTAGAGCCAGAGATCC TGGTGGATGGACCTCATGACATCAACAAGTGTGCTGAGGTGACCGAGCGCGTTCTTG[C/T] AGCATGCTACAAGGCTCTAAATGATCACCATGTTCTGCTTGAGGGCACTCTGTTGAAGCCC AACATGGTCACCCCTGGTTCAGAGTCTAAGAAGGTCACCCCAGAGGTGATTGCTCAATACA CTGTTACAGCTTTGCAGCGAACTGTTCCCGCTGCTGTTCCGGCCATTGTCTTCTTGTC
>Gm03_4 2243699	TGAAGCCCCAACATGGTCACCCCCTGGTTCAGAGTCTAAGAAGGTCACCCCAGAGGTGATTG CTCAATACACTGTTACAGCTTTTGACGCGAACTGTTCCCGCTGCTGTTCCGGCCATTGTCTTC TTGTCTGGTGGGCGAGGAGGAGGCAACCCCTCAACCTCAACGCCCATGAACAAG[G/T] CCCAGGGAAAGAAGCCGTGGTCCCTTTCTTTCTCTTTTGAAGGGCACTTCAGCAAAGCAC TCTCAAGGCATGGGGTGGGAAAGATGAAAACATTAAGAAGGCTCAGGATGCTTTATTTGCC AGGTGCAATGCAAACCTCACATGCAACTTTGGGAACCTACAAAGGTGATGCTACCCTTG
>Gm03_4 2338317	CTTATATATTTTATATATACATAATCAGGACTATTTAAGAGGAAACATAAGATAACATGCTGT ATTTGTGCTACAAAATAAAAGCAAAAGGTTACTTGAGAATCTCAATCCACTTGTCTGCAATG GCCATGTGCAGGCGATAATCAAAGCCAACCCACCATCTTGCGTAGGAAGGCAG[T/G]ATG TTGGCATTCCACTCACCTGGAAATGGAAAGAAAATTGAATTGTCAACTACCAGTTTTCATAC AGTCAAATAGAGGGTTCATCATCATATTGTCTTTTTGTTCACTCAACTCCTAAAACAGTTTG AATAAAATGTTAACAGAAGTACATTAATTCAAACCTTAAGAGGACAACATA
>Gm03_4 2672131	CCCCAAAACAAAAAAGACAATTCGACATCAATACGCGCAATGTAGAAGAACATGAAAA ATTAATAAATAAAAAATAAAATTAATAAATTAAGTGGAAAGTGAGAGAAACACTAATTCT TCAAATCCGGTGGTTGCCCTGATCTGCTCATTGTGAGGAGAAGAGATCACAGATG[G/A]CAA CGAAGGAAGAAAATTTGGTTTCGAAGTGGCAAAACCTCAACTACGCTCACTCAAGACCCTA ACATCGGAGGCTATACTCTTTATCGTTTAAAGCCAGGCCCAGTATACGAAACGATTGCTCAA AGGCACATGGATCTGATAGCATTAGTAAACAGCCAGGACCAAAATATGGGAAAG
>Gm03_4 4043363	TCTGCAATTTCTTCTCTGTCACTCGACTTCTTTACATCAATATAACCAGCTACTACGACAAGG TTGACCAGTTCCCTCAATTTGCTGTTCTTACCAAAGCATTCTACCAATCCAGTTATGTTTTTC ACTCTGTCTAGCCTTGCCATGGAGAAAATTATGGGCTTTGACTTGTCTTTCAA[A/G]GAACCA CTGCCACAAAAGAGACGAAGATAAACTCATTAACTGTCATAGTTGCTTTCAACTGACAAATG CAGCATGCATCTTAACAGCTCTTACAAACAGCTTTAGATATTTCTATTCTTATCAGGACTATG CATGTGGAGTCTGGGTGGGTTTTTAAACAGGACATCAGTACTGGAAGGA
>Gm03_4 4509128	AATCAAGAGATTATTAGTGACAGAGCCCATAGTCATAGAGAACACATGCATTTGTACCTGGA AGAACGTTCTGGGCTCTGATTTGGGGAAGAATTGTTCCATTTGCTCTGCGTCTTAGAGCCA CATTGCGCTGAGGAGAATCTCCAATATGCTTGCGGATTAAGCTGGAACATGCAAAGG[T/C]C ATTTCACTGTTGCTTGAGCTGAAAACCTCACAACACCAAACACGTGCGCTCCGCGCATCGCAG CCATTACTTCTCCTCCTCTGAATCACTCGGTGCAAGCAATTAGCAAACGCCGCTTTTCAAGT TTCAACCCAAACAAATTTCAATTTATTTACTTACTTTTGGGTAATAATATTTTAAAT
>Gm03_4 4562570	TGGAATAAGAGTTAATAGCATATCCCCTTTTGGAGTTGCAACACCTCTTGATGCAAAGCTT TCAACTTTGAACCTGAGCAAGTTGAAGCTAATAGCTGCTCACAGGCTAATCTAAAAGGTGTT GTGTTGAAGGCTAGGCATATAGCTGAAGCAGCTTTGTTTCTGCTTCTGATGATG[G/A]AGT TTACATCAGTGGTCACAACCTTGGTGTGGATGGTGGCTTCTCTATGTTTAATGAAGTTATT CCTTCACACCAGCTTAATTACATGTAGAGCCAAAAAATATAGTTTTGTGGCTGATTGGCA TTAATTTCTTTATCCTCATCAAGTAGAGAACATATATGATGTCATGCAGTGCG
>Gm03_4 4581602	CCATTCCAAAGACAGTAACGTAAACAAGAAAGGTAAGGTAAGTCTGTAGATAGTAGGGGG AGTAACTCCAGAGAGAAAATTAGCAAATAACTTTTTAAGCACTGTACATCAGCTACAGGCAAA TTATAAATCTTCAGAAGAGAAAGCGGAGGTGCTATCAAGGGGTGATTACCGCAGTG[C/T]JAG GAGGCAAAATTAATAATTGAGCAGCGGATTGATATGAATTCTGGTAGTCAACCTACGATGAA GCTTGGTGGGAAGACTGAAACATCAAACGACAGTTTATCTAATCAAAGTAAAGACGATGGG GGTGGTGGGAACAAGCAACGCAAGAAAACCTTCTAAGTTTCTGAGAGTACGCCTTGG
>Gm03_4 4581620	CGTAAACAAGAAAGGTAAGGTAAGTCTGTAGATAGTAGGGGGAGTAACTCCAGAGAGAAA TTAGCAAATAACTTTTTAAGCACTGTACATCAGCTACAGGCAAAATATAAATCTTCAGAAGA GAAAGCGGAGGTGCTATCAAGGGGTGATTACCGCAGTGAAGGAGGCAAAATTAATAA[T/A]T GAGCAGCGGATTGATATGAATTTGGTATTAACCTACGATGAAGCTTGGTGGGAAGACTG AAACATCAAACGACAGTTTATCTAATCAAAGTAAAGACGATGGGGGTGGTGGGAACAAGCA ACGCAAGAAAACCTTCTAAGTTTCTGAGAGTACGCCTTGGTATGGTCTGTTTCAGC
>Gm03_4	CTTGCTTTTTGTTTCGATTGAGACTATATGTTATAGCGTATTAGTATAATATAAGCCTTTTTGT

4779159	CTTCCCACTATGATTGGTGGTTGTATCATTTTGTGTAGATTTTAGGTTGTTTTTCCCACT AGGAGTGTGAGGTTTCAGGTGGAAGTTCGTTTTTGTAGTGATGGTCACTTTT[A/C]TTCTGT ACAGTAGTAGTGTGTTGAATAAAATGAAGAGGCCGTGATTGGAGAAAGTAAAGTAGTGTG CATAGCTAGTGAAAATATGTAGTACCAATGTGAACGCCCAAAGAAAAAACACTGGTGAATTT TTGGATTCATTTTAGCATAAGTTTGGTTATGATTTTATTCATCGCAAAT
>Gm03_4 4916159	TTGTGACACTCACGTACTATAAGTTGTTATGAACTTTTAATTTGCGTTGTAACATTTCC TATTTGTTGCTAACACATTATATATTGCTAGGTTTTCGTAAATGTTGTCCATGTTGGCGCTGG GCCTATTTCTCAATCTGATCTAGACTTAGCACAAGCTTGTGGTGCCTGCATAG[G/T]TGGATT CAATGTAAAAAGTCCACCTACTGCTCTTAGTCAGGAAGCAGCTCGTGCCGGTATCAAGGTA TAACTTTTTAAAGCTTATGCTCATATTTAACTACATTGTGGTTATGCTTAAGATTCTGTGAG ATGAAACTCTGACTTCCTTTTAAACATCCATTTGGTTTGGTTATATAGATA
>Gm03_4 5034077	AAACATTTTTATATGTTAACACCATTTTCTTTCCCGGATGTCACACAGATTACATTTGGTGC CAAGAAGCAAAATGTTGCAAGAAGCAAAATGTTCCGATCTCCTCTATATTTAGTAACGATA GCGATGAAGACTAATGATGGAACATTACTCGTGAAGTGACATATCAGCGTGATG[T/A]TGAA TCTCCTGTGTGGTTTGGTGTGGTTTGGTTTGTATTAGCTGTTTTATTTGATAATTGTTGCCAT CCTCGTTCAAGAGGAAGAAAACATTTGGGATTATATGATAGTTTGGAGTTTGGTAGACCAA AAATGTTACCTGCTTATCTGTGAGGTAATGTTATGTTTGAAGGATTTTAT
>Gm03_4 5035493	CTTCTGGTACCATTTTGAATTGGTTTTGCCTTAGCAGATAATGACATTAGTAATTTTCCAT TGGCACAAGTCATTGACTGGAGAAAGCAGGACAAGTATTGGCATCAGCATGGCACATTACA AGTATCCTAATTTTGAAGGGACAGAAAAGGCATAATATTCTGCCAACACGGAA[G/T]AAA ATAAAGAGTGAAGACGATGGTTTTTTCATATAAAAAAGCACACAACACAACACTTCCAAGTTC CAAGTGGATACTTCTACGTCTTAAATGGATAAATCCATTCGTGATTCCATAAAGTGAATCCC AGCTACAATTGTTATAACAGCTTATTTACAACACGCTTAAACAATTTTCAAAG
>Gm03_4 5035574	GGAGAAAGCAGGACAAGTATTGGCATCAGCATGGCACATTACAAGTATCCTAATTTGAAA GGGACAGAAAAGGCATAATATTCTGCCAACACGGAATAAAATAAAGAGTGAAGACGATGG TTTTTTCATATAAAAAAGCACACAACACAACACTTCCAAGTTCCAAGTGGATACTTCT[T/C]CG TCTTAAATGGATAAATCCATTCGTGATTCCATAAAGTGAATCCCAGCTACAATTGTTATAACA GCTTATTTACAACACGCTTAAACAATTTTCAAAGAAAATCCTCCAGAAAGCCTAATTAGACA GCTTCGGCAAAACCAACTTGTAGATTTCCGTAGTCAAAGACGGTGTGATATGC
>Gm03_4 5090113	AGCAATGGCTTCATGGAGCTATGCGTCACCATGGATACCTTACGTGGGAAGCACGTATCAA GATTCTCCTTGGCACAGCCAAAGCGTGAGTGTCTCCATCTGTGTTGCTTCTATTTTTTCCCT AGTCCTCATAAACCAATTTACATGTTTCAGGCTAGCGTATTTGCATGAAGCAATTG[A/G]ACC AAAGGTGGTGCACCGTGATGTAAAGTCAAGCAACATATTAATTGATGATGATTTCAATGCCA AGGTTTCTGATTTTGGCCTGGCCAAATTATTGGGTGCTGGGAAGAGTTATGTCACAACAAG AGTTATGGGAACTTTTGGGTCAAGTATTTTGTGCTCTGCTGAAGTTTTCCCATG
>Gm03_4 5098611	AGTGATAAAACCACTAGCTATGAAAAGCGAAATTGAAGTCACAAGAAGGAAAAGAAAT CACTTGGATATGATCTTTACATCAATTACAACCTTGCATACTGCAACACCTCCTCCTCATTC GGAGCTCCACTATACTCAGCATAGTTTCCAAGCTCTGATGCATAGCCTAAATCTC[G/T]AAC CTAGTAAACAAAAGGTATATTAACCTACCGAAAAGACCTACTAAAAGAAAACCTTCTAAAAAT ATCTGTTTGAGGTGTCCCTTTATTCATATTTTAGCCAGAATTAATATATCCACAAAACATG AAAACCTACCGTATCAGCATAAATTACACTTGCACCACCTCCTGCTACCATT
>Gm03_4 5105553	AAAACGTAGGTTGCAGAGCTGGGATGTTTCAATTATTTTCTTCTCATCTTTGTTGTACCATG GATCTGGAACAGAAAGACCTAGGCCGCATATCTAGAAGCAAACATGGGGCGACTTTGAA GATAGAAGGATGCGACCATGGACTCCATTATCGGTGGTGAAGAGCGGGAGCAAGCGG[A/G] JTAAGAAAGAGAGCATCGTGAACGTGGTTCAGTCTTGGCTCGTGTGATCTATAATGGCTTAG CCTTGACGGCGGCGGTGTGTTTTCCGTCTCCTCCACTTTTCTATCTTCTGACGGTGGC GACGATTGCGTTTAATTAGTTTTTTTTTAAAGTAGGGTTAAAATGGCCTTCTCATAT
>Gm03_4 5105650	GAAGCAAACATGGGGCGACTTTGAAGATAGAAGGATGCGACCATGGACTCCATTATCGGT GGTGAAGAGCGGGAGCAAGCGGATAAGAAAGAGAGCATCGTGAACGTGGTTTCACTTTGG CTCGTGTGATCTATAATGGCTTAGCCTTGACGGCGGCGGTGTGTTTTGCCGTCTCCTC[G/ A]ACTTTTCTATCTTCTGACGGTGGCGACGATTGCGTTTAAATTAGTTTTTTTTTAAAGTAG GGTTAAAATGGCCTTCTCATATATAATTTTATTTTGATATATTTTTTAAACCAACTACTTCAT TTTAATATATTTTTTAAACCAATTACTATCTTATAATATGAACCGTCCAATTATT
>Gm03_4 5107141	CTGGACCTGCTGCAGTTAGACCCGGAGCACCCAGTTTTAGACCTGCACCACCAGGGAGGT TCAATGATCCATCAGTACCTCCTCCAGTGGGACCTTCCAATGCCCCACCGGCTGCTGGACC TTTCCAGCAGTTTCCGCCACCGCCTTTTCCCCCAACAATGCAGCCACGTGGTCTCCG[C/T] AGCCAATGTTACCGCCATCAATTCAATCTCCACCTAGTCAAGCACTGCCTTTTCCACGTCT

	CTTCCGGCTCAACCACAGATGCCTTCTGTTCCAATGGGCTCTCCCCCGCCGCCGCCACAAA GTGCTGCTCCTGCACATTTAGGTTCAAATTTTCTCCTCCGCCACCTACAATTCAGC
>Gm03_4 5107353	CCTAGTCAAGCACTGCCTTTTCCCACGTCTCTTCCGGCTCAACCACAGATGCCTTCTGTTT CAATGGGCTCTCCCCCGCCGCCGCCACAAAAGTGCTGCTCCTGCACATTTAGGTTCAAATTT TCCTCCTCCGCCACCTACAATTCAGCCATCTTTTCTGGGTACCCGAGTAAGCATGC[C/T]A GTCCTGAAATGCAAGCTCCACCTATGCACTCCTCGTTCCCTGCTAATCAAGGAAATTTTGG GCCTGTCCCGCCTGCGGCTTCTTCTCCTTTTTTGTCTCACCCAGGAGGTTATGTTCCATCAC CCCCAATGGCTCCCCACTAGGTATCCAGCCAATGCAGCAGCCAGGATCTGTGCC
>Gm03_4 5114910	CATAATCACTTATCAACTTTGTATATTGATTCCCTTTTGGCAGCTTCCCTGGCTAAATAAGCA AGTCAGGTCTGCTATTTGTCTCACAAGTGACGACCATTTCTTTCTGCATTGGAGATATATTA AATCTGTCTTTGGGGGTGCTTGGAGAAGCATTCTTGACCTGGCTAATTGATCG[T/C]GACT TGTGAGAAGCATGCTTTTGAAGATTGTTGATTATCCTGGGATGTATCCGTTAGGTGAACT CTACCACAGAGATTGAATTTAAACGTGTTTTCATGTTTTTATATAGAGGGAAGAAAGAAAC ACTTTTTTGGTTACTTTACCTTGAGATTAGTTGTCACTTTCTGTTTTATA
>Gm03_4 5300266	TCTTTCACCATACGATAAGGTCATCAAAGCCTACCAGAAAAAGGGGGAAGGGGGGATATGT ACAATCTCAAACTACATGGAGAACATAACTAGGAATCTGACTGCCCCATGTTATGTCAGTA AGAGATTGTGTATGTTGTGCATTCAACTGTGCCTGGCGTGCAATCTCCAGTTGGTC[A/C]AA TCTTTGACTGAAGTAGTACAGCCATCGCTGAAGCCTTATTCTGATGTTGCGGTCTGTTTTTA AATAATACCCAAATTCAAATCTAATAACACAACACAATCAATAAATGAATAAGTGCACATAT TTCATACCTCTCATTCTGACAAGTTGCAGTAACCTCTGTAGGAATATGAACTA
>Gm03_4 5329182	AAACGTTAAAAAGAGTGAGGTAATCCAACAAGAAGAGGGGAAAGCAAGAGTAGTGACAGA GAGAGGCGAGGATCGTTTACATAAAACATCGACAGAGATCGACGAGGAAAAGCTAAGCCT ACAATGTGGAGCTCCAAAACGACATTGATGTTTGGCGTCGCAGCCATTGCTCTGATGA[T/ C]GTTTCTCTCATCTGCTTCAGCGGACGACGTCGTTGCACTCACAGAGGAAACCTTCGAAAA CGAGGTTGGCAAGGATCGTGCCGCTCTCGTCGAGTTTTACGCTCCCTGGTACGCGCGAAT CGTGATTCTCTCATCTCACGCGCGTCCATTTCATTATTTCCATTACTATGAATTATGAAA
>Gm03_4 5336647	AACCCGAACCAAGTCAACCCGACGAACCCGAGGCGGAGAACTCAAAATCGCAACAAG AGTTTTTTTCCAAGGAAGACCAAGAGGGTGAGTGTGAGTAAGTGGAAGCGGCGGATATAGT GAAGTGCATTGGGCTGGCAGGAGCACTGTAGAAGAAGCCGGCTCCGGCGGCGGAGATG[A /G]ATTCCCGGCCGGGACTGGAAGGGGCGCTGACGTAGGGAGTGGAGCAGGTGCTGTGCA TGTCTTCAAGAGGGGAGTGTCTGGGTTTTCGGTTACGCCATAATTGTTTCGGGGTTTGGAG CTCCATTATTCTGCTCTCTGTCCCAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAAA AG
>Gm03_4 5392239	CACTATTAATTGATTTTTCAAGGGTAACCTACTGAGTATTTTTGTGCTCTTGTGAATTTCTGT AGGATGGAGGACCTAATTTCCATTTTATTCCAATACGTTTCAAGTCTGAGAAGCCTGGGG GAGATGTGTTATTTGTAGCTCACAATGCTCGCTGTTTGTATGTGCCCTTCATCAT[C/G]AGTG AGTTACGGCGTTATTCCATAAAGATTCTCCTAATTGGTTGTTTTTGGACACCTTCCTTTAG CGCGGGAACCTAATGAAGTCCGGAGGTCTCTTTCTCTCTCCTCTTCTCTTAACACATATGCAT GCACGCACACGCACACACACACAAATACTTATTGTTTCCATGTGAAAACCT
>Gm03_4 5416367	GGTTGCTTGTTATAGATGTTGATGCTGCCCTACCGTTCAATATTCCATTATCCATAGGTGGA CTGGTTCCCCACTTACCTTCTGATTCACTAGGCTCTATCACTTTCACGGACCCATCTGTCAA TCCAACAGCAAATTGGTTGGGTTCCAGTGGATGAGCCGCAATGGCAACAGGATAC[G/A]CA GATTGATTCTGCAATATAGTATACCACAATATGGATCACCAAATGGAAGTAAGACTTCACG ATACTACATCCTTAAATAAATGCTCAATGTTTAAATACAATAGAGAGTAAATTTGGGTCTGATC TGTTGGGCAGAACTTACCCGCTTAAAGCTGCTGGAGGTGAAAAGTATGTGGA
>Gm03_4 5471080	ACATTGAGACGCTGGACCTGTGATGTGTCCGCGGATCGAGGACGGAGCTGTGTGCGTTG TGCTGAGTCAGGGATCGGCGAGTTGGACTCGGGGACTGAGGAGACTCGTGTGATCGC GCCACCGGTTGGGGCATGTGGGCTTGAGATGCTGATTGCGGCGGTGCCATGTTGGAG [C/T]CCGTGGATGTGTCCATTGTTGGGGGTATGGCGACAGAGAGGCTGCGGCGCTATCGT GCGCCGCGAGGTTGAGGGAACCTCAACATGGATAAGTGTGTTGGGAGTTACTGATATTGGGT GGCCAAGATTGCTGTGCGGTGTGGGAAATTGGAGAGGCTGAGTTTGAAGTGGTGTGTTGA GA
>Gm03_4 5493120	GAAATCTCAACCTGTAAGTACCTTTTTTCCAATGTGAATTGCAGGATTGTTTCAATATAGCTTA TTCTATTACTGTGCATTATATCTATACAGAAATTGGGAACCTCACAGCACCATTGGACTGGA GTTCAAGTCAAAGTTCCCGTGAAGTTCATTACAGGTGATTTGGATGCAGTATAC[C/T]CTTCA CTAGGGATGAAGAACTATATAGAGAGCGGCGCTTTCAAGAAAGATGTGCCATGTTTGGAGG AAGTGGTTGTGCAGGAAGGGGTGGCTCACTTCAACAACCAAGAAGCTGCAGAGGATGTCA

	CTAATCACATTTATGATTTTATCAACAAGTTCTGATGCTATCAGCTTAGTTAAT
>Gm03_4 5503654	CTAAATACACAAATAGAAAATGTTTTGAAAAGGAGCAGGGAGTGCAACATTATAAAGAGTG GCCACTCAATAGTCAATAAGGGGAATCAAAAAATCAAATGCTAGTTTTTTGTAATACACGTC AATGTA AACATGGGAGCCACGATATGTTATCAGAACAGAAATATCATAAAAACTG[T/C]CAC CGACGCCTAGAGCAAACAGTTTAAATTTTCAAGAGTGAACAATTGTAAAAAATAAGAACAA AGTTATATTATTTCTTAAACTCTAAATCCTCTGCTGTTCTCCTTCTCCTAGCCCTCTCGA ACTTTCTTCCCTTTGCTCGTACATAGGGCTTGGTATGGCTGTGTGGCACACC
>Gm03_4 5503673	ATGTTTTGAAAAGGAGCAGGGAGTGCAACATTATAAAGAGTGGCCACTCAATAGTCAATAA GGGGAATCAAAAAATCAAATGCTAGTTTTTTGTAATACACGTCATGTAAACATGGGAGCC ACGATATGTTATCAGAACAGAAATATCATAAAAACTGTCACCGACGCCTAGAGCAA[G/A]CA GTTTAAATTTTCAAGAGTGAACAATTGTAAAAAATAAGAACAAAGTTATATTATTTCTTAA ACTCTAAATCCTCTGCTGTTCTCCTTCTCCTAGCCCTCTCGAACTTTCTTCCCTTTGCTCGT ACATAGGGCTTGGTATGGCTGTGTGGCACACCAGGAGCAGGACCAAGTGT
>Gm03_4 5516926	AGGAGGAATGAACAATACCAAAACAAACGCAGAATGGCTTATTAATTATAAATATACATTA CATGATAAGGAGTAAAGATACACCGCTACTCTACCCTTACACTTTCTACTTACGATTTGGCA ACCACTACGCAGCTTACTTAAGTAAAGACTTAAAGAAGCTAAACCAAAAGCTACC[T/C]CGC ATGAACCTTGGCAGAACTTCCTTGCAGACATTACAGACATTATACATGCACACAAAGACTAAAC AAAAATGAATACAGCAAAGGCCAAAAAAGCTGCTACATCCTCAGCTTTTCAAGTACGAGAAA ATGCGTCACACATCGAAAGGAGAAGAAGGGCCTGATCTGTTCTGCCCTCTCTTC
>Gm03_4 5516951	AAACGCAGAATGGCTTATTAATTATAAATATACATTACATGATAAGGAGTAAAGATACACCG CTACTCTACCCTTACACTTTCTACTTACGATTTGGCAACCACTACGCAGCTTACTTAAGTAAA GACTTAAAGAAGCTAAACCAAAAGCTACCTCGCATGAACCTTGGCAGAACTTCCT[A/C]GCAG ACATTCAGACATTATACATGCACACAAAGACTAAACAAAAATGAATACAGCAAAGGCCAAAA AAGCTGCTACATCCTCAGCTTTTCAAGTACGAGAAAATGCGTCACACATCGAAAGGAGAAGA AGGGCCTGATCTGTTCTGCCCTCTCTTCTCGTACGCATCAGATCCCGACCTTC
>Gm03_4 5517710	ACTTTTACTGGACCGTGTGTGTAATTAAGTAAAGCATAGTATTTTTACTGTACAGTAGAAA AGAAATAACGACCTTACCGGAAAACGGAGAAATCGGAGGAGTAGAACCAGGAGAGAAAC CGGAGGCGATCCACTCTGGTACCCTGGCGGTTTACGATCATGATACTACGCGTCA[C/T]T CTAGTCGCTTCTCCGACGATTTCGTCACCGTACGATCTCACGCTCCCGCCCTCCGCTTCTG ATCAATAATCAATAAATAAATATGCCTAATCGTTAGCATATTTTGATGTTTTAAACGGAGAA AAATATCTATGAGCGTTTCAAGTGAATAGATATCTCCAAATTAAGTGGTTAG
>Gm04_5 9457	ATCACCAAAAAACACATTTAGCCAAAAGCAATAGGAGTTTTTGAAGTAAAAAATGTTAAAGA GAGAGAGAGAGAGGACCTGAGAGAGAGAGGACCTGAGAGAGAGAGAGAAAGTCTAGAACA TGATCGGTGGAGGTGACTGTGACGGCGAGGATGAGCGAGGCGAGTGAACGGAAGGAAA[T /C]TTTAGAGCGAGTTGACCTCCGTCACTGCTCCTCCCGTCCCATCAGTACTTCTACTTAAAT ATCCAATCAACATTATTTACCTATCTCCTAGTCCCAAATCAACAAGTTTTGGGTGGAAAACCT TAAGAAGATTCTTATAAAATAAAAGTCAAATTTTAAATTAAGTATAGTTTTTGTGTC
>Gm04_1 36286	AACAAATGGAAGAAGTGGAAGATGGTGTAGCAGAAGAATCCATGGGTGCTGGAGCAGAAA TAATGGAAGGTGTTGACGTAAAGCCCAAGGAGGAATCCGATCTAGACGCTTCCCATGA GGAACCTTGAGCGTCTGCAGGAACAGCTTTTCAAGACAGTATAGCTGTAAGATCTAGGCGC[T/C] AAACAGAAAGAAAAGCTAAAAAGAGTCAGAGCAACAGAGAAGGCTACCACAAATTTTACATC ATTTAAGCCTGGCTCCTCCAGCAGGAGAAAGCGTGTTTCTATGCAAGAAGTAGATTATTCA GACCCACTTCGTTACTTAAGAACAACAACCTAGCGCTTCTAGGCTTCTTACTCCAACCG
>Gm04_6 33262	GAAACCATTATAATTACAGCAAGAAGTATATTGCAGAATTAATGCTAAATCTTAAGAAAC ATAGAAGGAATTCCATACCCAGGGAGGCCACAGTTGTGGCAAATTGCGACATTAGGGCATT CTCTTGACATAATGACCAGGTCTCTTGACAGTTCTTGACACAGATTGTCACGGCTGAC[G/C]AAA AAACCTAATTAGTGTTGCCATTGTAAGAAATGCATGCTCATGTTCCAAAAGTCAGATGGCTT TGAGTAAATGCACCCGCTCTGTTTGAGAGGCAAAGCACATCACTTAAGTTTCAAACCTAATCA TAGCAGCAGCCAGAGTAAGAAGATTGCAAAACCAAGAACAGAAAATAAATTG
>Gm04_9 88779	TTTTTCCAAGAAAACCTCCTCAACATCTCCAAATTTCTGACCAGAGAAGAAGCTGATCAGA TTCTTTTCTCATCAAAACACCTACCATCGCTTCTCAAATTTCTTCTCCATCCCAAAACACTCTC CCCAAGCCAAGGCAATGAAATATACCCTCAACAATGTGAATTTGAATCCATGG[T/G]AGAG AGACCAAGTTCTGTGCCACTTCCCTGGAGTCTNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
>Gm04_3 9101834	GTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT GGAAATGAAAGAACAAATAACAAACCTTTGCCAGCTCCCATAGTGTGACGGGCGGTGTGTAT

	AAGGCCCGGGAATGAATTCACCGCCGTATGGCTGACCGGCGATTACTAGCAATTCC[A/C]A CTTCATGCAGGCGAGTTGCAGCCTACAATCTGAACTGAGGACGGGTTTATGGAGTTAGGTC ACCCTCGGGGATCGCGATCCTTTGTCCCTGATGTGCCATTATTTCTCCTATTTCTTAACC CCTTTTGTACCATTTTAAGTACTGATTAATCTTAATTGTCAAATTAATTAGACAGT
>Gm04_4 1571642	TTCTTTTCGTTCTTCCACAGCCGGTAGGTACTGTTTTCACTCTCATTGTTTGAATTCTTCTCT GTTTGTTTTAATGGCACTGCCACACCGTATCATTACCCTCTATCAAACCTAATTTGCCCCCT TCTCTTTATGAATGAATTCATTGAATATACGAACTTAGGGTTTTTAAATAACTG[G/T]AGATT GGCTTTGTCCATGGCCACCGAACCAAAAGCAGCCACTGAGGACGTCAAGATTGATCTCTTC GAAGACGATGACGAATTTGAAGAGTTTGAATCAACGAAGGTAAATCTTGTCTTTCCGCAC GCAATCTCGTTGCCCTCTGTTCAATTTTTGTTTTGTTATTTTTCCCAAT
>Gm04_4 4751336	CCGTTAAACTCAACTGCAACTGCAGCAGCGAAACCAGCAAAACGTGTTCTTCCATGCTCA ACGTAGCAGAACGCGAAGAAGCACTACCAGATTCCACTCATCCAAACGCTGCTTTAGTTCT TCCGCCTTTCCAAATGCGTCCGCTTTCCGCCGCGCGCGTGTGCCAGGTCGCCAGC[A/G] JTCGACTTCGAATCCGCCGCCAAAGCCGGAGAGGATCGTCTCTCTAAGGTATCATCGCTAC TACTACTATTTCTCGTGTTTTTCCCAAATTAGCCCTCGTTCTGTAGCTACTAGTACAAGCTC GTTTGGTTCCGTTTTGGTTTTAGGTGCCGGTGCCTAATATAAGGAACCTCTGCATCA
>Gm04_4 4761104	CTCTTTTAATTTCGCTTATTTTACTTTTCATCTTCACTACATTTAACTGGTTTAAATTTGTCGGC GATTGAGCTCATCCAGCATCAACAACCTCAACTTTACCCACACTTATCTACCATTCTTCAAAT TCGAAGCCTCCTAGCTCGGAAGTGGAGTGTTCGGTTACAACACTGCTATCGT[G/A]AAGGCA ATGCCCCAGCGGACTGGTTAGCTAAGCATGGTGCTTCTAACGCACTTTTTCTCTCCGTCCA GCCTCACCTTGCTGCTTTTGGCTGATTCTCTGGAGCTTCCATTGCTAGAACCTAGTTCTGT TTTCTGCTTGCTGTCTTGTATTTCTTACATGATAAAAAAAAAAACTCGTTT
>Gm04_4 4787477	AAAAGATAGAACCCCAACAAAAATATGTATAGTTATATATAGAGAGAGAGAGAGGGCCTTG GTTGGTGGTGCAAGGCATAAAGAGGAGAATTGGTGGGGATAGGCGAAGGAGGTGGCGAC TGAGTAGTGAATAGGGTTTTTGAATTGATTGATAGCAATAATTGGTGTAGCGAAGCTTG[G/A] GAATTGAGATTTATGGGTCGCAATCGCAGTCGCAGTCGCAGTGTGTACGCTCTCCTTCCA ATTCCCGCAGAAGAAGGTAAGTCTCCATCCCCTGTTTCTCATCGCCACTCGAGAAGTCCAG GCGCCGCTCTCCTTCCCACAAGCGCCGTCGAAGGCACAGAACCTCTTCTCCCCCTCT
>Gm04_4 4787569	GGTGGGGATAGGCGAAGGAGGTGGCGACTGAGTAGTGAATAGGGTTTTTGAATTGATTGAT AGCAATAATTGGTGTAGCGAAGCTTGAGAATTGAGATTTATGGGTCGCAATCGCAGTCGC AGTCGCAGTGTGTACGCTCTCCTTCCAATTCCCGCAGAAGAAGGTAAGTCTCCATCCC[T/C] TGTTTCTCATCGCCACTCGAGAAGTCCAGGCGCCGCTCTCCTTCCCACAAGCGCCGTCGA AGGCACAGAACCTCTTCTCCCCCTCTCTTTCACGCAGCCCCACTCCCAAACCCAAGAAAG ATCAAAACAAAAGGTAACCAATCATAACCATTCTTTATTTGCCTGGATTTTATCTTA
>Gm04_4 5033967	ATGTCATACAAAAAGTGTGCAGAAATAACAATAGAATAGTTAGTGGATTGTTCAAAGATAGA CATTGACTGTAAGATGAAAAGTTGAACTTTCCAAATGAAAGAAAGAAAGGAAGAGAGT ATTATTCAAATACCTGATCTTGGGGGAACCTTCACTTGCACAGGAGAGCAGTGATCA[C/T]GA TTGTCACCGACTCCAACCTGTCCAAACTACAATTGAAGGCAAAATAGGTAAGAATTTTCCAA ACACATGTCAATAGTAAAGACTTGTGATTGTACAGAAAGAGAATAAAGTCTTAATACCAAT TTTTTTTTTATATGAAGTAGAATATATTTAGTTCTCTGCACACATGAATGTGCA
>Gm04_4 5034787	AAATGCCAATGCATCCAAATAATACAAGAAATTGCTAGCTAGACACCAACTCAGCATAAAAT TAAAGGACTACCTGAGAGATTAACCTTATCACTCAAGGCTTGAAGCTTCTGAGGCACAAGAG AATCCTCAAAATTTCCATGTCTAGCTGGCCATATTTGCTCCACCCGTATGTGTAT[G/A]TGC CACCAGAAGATGAAACAGAAATTGTATGCCGCCAACCACAAGCAACCATGACCATCTTGTC ACACTGCAACATTAACAAGCAAGAATTAGGCAAGGGGAGGAGGGAGAGCAAAAAATCAAAA GTTCTCCATCCATTACTATGCATGTACACCAATCACACATGAATTTTAACAACAT
>Gm04_4 5061078	TTTTCTAGAATCTTAATATTATTCTAATACTAATATAGTATTACCATGCAGATGTTGGCACT CATGGGAAATTCCAAGACCAAAATCAATTGTCCTAACTGGCCACTCCATTGGAGGAGCCACA GCCTCTCTATGTGCTCTTTGGCTACTATCTTACCTCCACCAGACATATTCCTCTA[G/A]TTCT GTGTACGTCTTGTGCATCACATTCGGTTCCCCCATGCTAGGGAATGGCTCTTCTCACGTG CCATTCTCAGAGAAAGATGGGGTGGTAACCTTCTGTACGTGGTCTCAAAGCATGACATAAT GCCAAGGCTTCTCTTTGCTCCCATACACCTTACACTGCTCAAATAAAGTTCTCT
>Gm04_4 5061509	AGAAAGAGCTGTTCAACTTTGTGATGAGTCACTTGGATGCAGCCACACAAGATGAAGAAGG GTCTGCACCTGTTTTGTTTCACCCATTTGGGAGCTACCTTTTTGTTTCATCAGATGGAGCAG TGTGTGTGGATTGTGCAACTTCTGTTATAAAGATGTTGCATTTGATGTTTGCATCA[A/T]TTTC CCCAGCTTGTAGTATTGAGGACCATCTAAAGTATGGAGACTATGTTAAGAATTTGTCATTAC AATTCTTGAACCAGAACAAATCCGTGCAGGGGAATATTCCTGATTCAAGCTATGAAGCAGG

	ACTTGAATTGTCGGTTCAATCTTCAGGTTTGGGTAACCAGGTAAATGCACTAT
>Gm04_4 5090419	CCAACCCCCCTCCCCCAATAAAAAATAGGAAAAGAAATTAGCTAACCCTAATGACTGATGAT GTAAACAGTTATGGAGGAACCGTTGGCAGCTTGAAGAACATTGCGACTAGAAACATGTCAA TTGCTTTGCAAACCAAGGCATACAATCTGACTGTTTTTGCCTGTATTCTTACAAAT[A/G]CTG TTGCAAAATTGGTGCACTTACTGGAGTTTGATTGGTTTTACTATTTGCAGAGACATTGAATTT GTACAGATTTGCATGTATAATAATACATCAGTTTTTAACAAGAATTTGACTGAGAAAATCACA CGCCCCGTCCCTCTTGCAACTACTTTCACATTATTTACGTTAGATACTTCC
>Gm04_4 5152573	ATATTGGTTTCAGAATTAGCTGAAGTACTATCAAGTTTGTCAATATCATGCCAAACACCAGT CCTTCTTATCTCCATCTATCAAAGGTAAAAGTTACCATAAAAGCAGACAACATTGAATTTTCA AGTGCTATCAAGTACCTGAAAGTTGTGCAATAAGCGCTCAGTAATGTTATCATC[C/A]TAGTA CTTCATCTCATCCCATGGTCCATCTCCAACACCAACCAAAATAATCGAGAGAGGATAATGAC TGATAATCAGAATAGCTCCACATTTAAGACAGATACATAAAACAACAAAAGGTTACAAAGATA TCACCAAAAAAATAACAAGAGAAGTTACTCTACCTGGCAGCAATGATAGAA
>Gm04_4 5374514	TCAAAACATGGACCCTCACCGCCGCCACCGCCACCGCCGCCGACGTTGGTGCTGCTCCTT TGCCGTCCCTCCCTCCAGCCCCGACAACCTTACTAATCCTCCAAACCTCTCCCTCGCAAC TCCGAGCCTCACTCCAAACCTGCTTCTCCCTCCAAACTCCCCAAATCCCGCATCC[T/C] AGTCAAGATCGACCCTCGCCGGATCCTTTCCCGGGAAGAGTCTCCCCCATCGACTCCGA TCCCGTCGCCGGAGAACCTCTTCCCTTTCCCCCTCCCTCGCCGGAGCCGCCGCCGCCGCC ATCTCCGGCGCGGGCTACCGGCGAGAGGGACGTGAGGATGAGCCTGAGAGGAAAGAACG G
>Gm04_4 5380764	AAAAATAGATTGCACACCTCAATTCTGTTGTTTTCTTTCACAGCCAATGTGTCATCATCTTCA AAGAAACAATCAAGCAAAGATAGAATGTCATTGTCAACTTGATTATTTGGAGCTGAAGGATG TGCTGAGCAAGAGGTCATCAATTCTGGTACACAAGATACAGAGCCAACCTTGATT[G/A]CTTG GTGCTGAAGGATGGTCTGAGAATGAGGGCATCAATTCTGATGCAGATGACACAGAGCCAA CTTGACAGTTTGGTGCTGAAGGATGTGCCGAGCAAGATGGCATGAATTCTGAAACACGTGA AACAGAGCCAACCTGACTGTTTGGTGCTGAAGGATGTGCCGAGCAAGAGGGCATG
>Gm04_4 5380790	GTTGTTTTCTTTCACAGCCAATGTGTCATCATCTTCAAAGAAACAATCAAGCAAAGATAGAA TGTCATTGTCAACTTGATTATTTGGAGCTGAAGGATGTGCTGAGCAAGAGGTCATCAATTCT GGTACACAAGATACAGAGCCAACCTTGATTGCTTGGTGCTGAAGGATGGTCTGAGA[T/G]TGA GGGCATCAATTCTGATGCAGATGACACAGAGCCAACCTTGACAGTTTGGTGCTGAAGGATGT GCCGAGCAAGATGGCATGAATTCTGAAACACGTGAAACAGAGCCAACCTTGACTGTTTGGTG CTGAAGGATGTGCCGAGCAAGAGGGCATGAATTCTGAAACACATGAAACAGAGCC
>Gm04_4 5380834	AATCAAGCAAAGATAGAATGTCATTGTCAACTTGATTATTTGGAGCTGAAGGATGTGCTGAG CAAGAGGTCATCAATTCTGGTACACAAGATACAGAGCCAACCTTGATTGCTTGGTGCTGAAG GATGGTCTGAGAATGAGGGCATCAATTCTGATGCAGATGACACAGAGCCAACCTTGA[C/T]AG TTTGGTGCTGAAGGATGTGCCGAGCAAGATGGCATGAATTCTGAAACACGTGAAACAGAGC CAACTTGACTGTTTGGTGCTGAAGGATGTGCCGAGCAAGAGGGCATGAATTCTGAAACACA TGAACAGAGCCAACCTTGATTGTTTGGTGCTGAAGGATGTGGTTCCGAGCAAGAGG
>Gm04_4 5380885	GATGTGCTGAGCAAGAGGTCATCAATTCTGGTACACAAGATACAGAGCCAACCTTGATTGCT TGGTGCTGAAGGATGGTCTGAGAATGAGGGCATCAATTCTGATGCAGATGACACAGAGCC AACTTGACAGTTTGGTGCTGAAGGATGTGCCGAGCAAGATGGCATGAATTCTGAAACA[A/G] GTGAAACAGAGCCAACCTTGACTGTTTGGTGCTGAAGGATGTGCCGAGCAAGAGGGCATGA ATTCTGAAACACATGAAACAGAGCCAACCTTGATTGTTTGGTGCTGAAGGATGTGGTTCCGAG CAAGAGGGAATCAATTCTGATACACATGATACAGTGCCAACCTGGATTGCTAGGTGCTG
>Gm04_4 5381162	TGATGGCAAATGATGTGGTTCCGAGCAAGAGGGAATCAATTCTGATACACATGATACAGTG CCAACCTGGATTGCTAGGTGCTGAAGGATGTGTGCGAGCAAGATGGCATCAATTCTGATACAC ATGATACAGAGGTCAATCCAGCACATCCACTTGTAGAGGGATGGATGTAATTAGCAA[C/T]T GAAATATGGCTTTCACTAGGTTGTATAGGAACAGGTGCACAAACAGCAGCAACAGGGGCAG ATTGTACAGAATCAATTTCTCATCTGTTTCCCACTCTTTCTCCTCATATGGTTTACCATATT GTGCACCATTCCTAGGACCAGGACCTCCTTTTGGTACACCCTACAAATCACATA
>Gm04_4 5385128	CTAACTCAGCATTTGGGAGGCTAATGCAATTTCTCCAGGTAGCAGAGCTCCATTACACAAGA AACAGGGATGGAAAAATCAAAAAGACTCGGCTCAAGTAAATTTAGCCTCAGACTGTTGAAG TACTTTTGCAGTCAGAATCTTCTCACACTACTCCAGGAGAATAGACAATGCACAC[G/A]G CCACCTTTTAATTCTTACTTCTCAAAAAACTCCAAAGCCATAACACACAGTGAACCAACCA TCTTCAAATGACTTGTAAGGCAAAATACTAATTGATGAACAATCACCGTGCAAGCTTAGTC AAAAGGGAACAAGCAAACGATCACCATTGACAACTATTTACAATATCCATTCT
>Gm04_4	CCAGCTAGCATGAAGAGCAAGATTTGCCAATAGGAGCGAGGCAATCTCACCTTATCCACAC

5580521	GCCCACCACAACCTTAAATTACATCCATCATCCATAGCAACAAGCCAAAGAAGAGAAAAATAA TAATATGAATAACAACCCACAAAACCTTGGGTCTTGCAAATTAAGTTTCATTTTCCTC[G/C]CTTT CACTCTCGTAGACATCACAAACCTTCCCTCCATTTCATCCACGCATGGCTGCCATGAACCT AGCGTATTGGCCTGTAGCTATGCCATATCAGGTGCAGCGTGTTCGGAGCTCAATGGGAAG GTCACCTCCGTGGCCTATGTTGCATGCTCTGGCTACAAGTTGCCATTGATCAAA
>Gm04_4 5580672	GGTCTTGCAAATTAAGTTTCATTTTCCTCCCTTTCACTCTCGTAGACATCACAAACCTTCCCTC CATTTCATCCACGCATGGCTGCCATGAACCTCTAGCGTATTGGCCTGTAGCTATGCCATATC AGGTGCAGCGTGTTCGGAGCTCAATGGGAAGGTCACTTCCGTGGCCTATGTTGCA[G/T]GC TCTGGCTACAAGTTGCCATTGATCAAGCTGAGGCCAGAGTTCTTAAAGCCAAAGAATCTG AAGGAGGAAGGAGAGGTGCCCTTGTCTCTTGGCAGCTACCTTTTACCTCTGCAGCTTC CAATCTTCTGCCAATGCTGGGATCATTGAAGATTACCTTGAGAGGAGCAAGGCCA
>Gm04_4 5581792	TCTCATGCTGTTGTTCTTTTTCTTCTTTTTTTTTCTCCTTTTCGTTTCTGGGTTTTATCTAC GCAGAAAGTGCCATTCCCTAGTGACGACTTGGAGCTTGAATGTGAAGGAAAGGACAAATAC AAGTGCGGTTCTAATGTTTTCTGGAAATGGTGAAAGGAGGCAACATTTTATGC[G/A]TAAC GCGAACTTTTCGCTTGCAATGTACCTAACTGCTTGAATATATATCATATTTTTCTGCATTGA TTTATTATTATATTGTTGCTGTTGGATTTATTTGTTGAAAAAATGTGTGCATTAGGCGGCT ATTTACTCTAGATTGCTTGAATACATTTTGCGATATGTTTTGATTTT
>Gm04_4 5581810	TTTTCTTCTTTTTTTTTCTCCTTTTCGTTTCTGGGTTTTATCTACGCAGAAAGTGCCATTCC TtagTGACGACTTGGAGCTTGAATGTGAAGGAAAGGACAAATACAAGTGCGGTTCTAATGT TTTCTGGAAATGGTGAAAGGAGGCAACATTTTATGCCTAACTGCGAACTTTTCG[A/T]CTTGC AATGTACCTAACTGCTTGAATATATATCATATTTTTCTGCATTGATTTATTATTATATTGTTGC TGTTGGATTTTATTTGTTGAAAAAATGTGTGCATTAGGCGGCTATTTACTCTAGATTGCTTG AAATTACATTTTGCGATATGTTTTGATTCCCAAATTCATGTTAGCTT
>Gm04_4 5594453	AGGTAATAAAAAAAGTTAAACAGGAAACTTTTTCGCTTTTTCAATAATTGGTTGGAGACAAG AACTATTTTTGTGGAAGACATCTTAGACATGTATAGTTTAGTTTATTTCAATTTTTCTTAAC AGGAAACAATCATTAGTAATTACTCTCATAAAATTGGTGATGCCAGCAACCA[G/A]ATTTTG GTGTTGACTGCGATGAGCAGTAGTACATTAGGCCTCATTGAGGTTTATGACAATCATGGCC GTGCTTACCCTTGCAAAGGAAATGAACATGAGAAAATCTTAATCTTATAGATCCCTAGAAAA CCTTAATAAACTCATCTACAAAATCATAGAAATGGACTCATAACTTCGTAA
>Gm04_4 5597626	CACACCAAATGGGCGGAAAAAACAATTTTCATGCAGTGAACAAATGGGTACATGAAA AAATAAACTTTTTTAAAGGTATTGCGAAAAAACAACCGAGAGAGAGAGTACCCGAAG AAGAGCATCTGACGGTAAAGACGGAACCTTTGGCGGCAGCATGTGGCGTATTGGAA[C/T]T GAAGCGGTTGTTGGTGAGGTCATTGAGAGACAGAGCGAAGGACAAAGCTTTTGTCTCCATT GATGCCTCACAGAAGAAGCAGCGTTACAGATATGTCTGAGGGAACCACTCCAGTCGCAGTA CCTAGGTTTCCGCCACCTAGGTTTTATTTTGAATTTTCGCTTCATGGGGCTCTTT
>Gm05_3 7147034	GCCACAAAATTTTCAAAACAAAACATCTGAAGGATAAAAAATGCCTCGCGTACTAATTA ATTTTGAAGATTAAAGTGGAAGGAAATTGGCTACGTGTTATTACTATCAAAATCTTTCCAG CACAAAGATATTGCTTTTGTGCTCCCTTCTATTCTCAAGGTATCATCAACATA[T/C]CTATG GCATAATGGAGTTTAGGAACAACAACAATTTTCATCAATAATAAAAGCATGAGAAAACAAA GAAAGACATGATTTGGGCTCGTCAAAAAGAGTAGATATGGGTTGGACCTCAAAATTAGACC AATAAAAGAAAAGAAGAAGAAAAATCTTCAAAGAAATTAGGATACGTGATTTT
>Gm05_3 8244445	ATTTATGTCCGTGGAGCTGGAAGAAACCAATGCCAAAGTTGTTTATTTTCGATATTTATGGGA AAGTTTTGCATGTTGTGAATCTGCCAAGGGGTTAGGTACTTCTGTCTCTGCCATATAGAAG ATGTTTTCTGTTTTCAGTGTTGCAAAAGGAAACGATAGTTATGTTTTGCATCC[T/C]TTAAT TATCCACAAAACAAAACCAACGCTCTGTCTTAACTCCTTTGTTTTATTCCATATTAATGAA GTGCGGAGATGGCTGCTATTGTATTATATATGTCTCCAATTAATTTTTAATGTGAGGACTGA ATCACTTTCTGAACGCCTAGATCCTCATCTTAATGGTCCTTGAAAGTATA
>Gm05_3 8251772	TGCTAATATTTCTTTGATACCCTCTGTAAACCCAGATCTGTCTGGTAAGGTTGATTTGTGGC CAAATGAAGATGGTAAACTTGAGCAGCTTCACTCAGATGAAGCATATGAGATGATTCAAAAC CACTTGAGTCTCATTCTCGGTAACAAAGCTGGAGATTTAACATCAGTAGCTGAAA[G/C]CAG CAAATTCAGGGTAGGGCAGGTCTATGCAGCATCAGTGATGTATGGTTATTTTCTTAAGCGA GTTGACCAAAGGTTCCAGTTGGAGAAGACAATGAAAGTTCTTCAAATGCAACAGAAGAAG AGAATGGTGTTTCATCGAAATACGATGGACAATGCAAGACCCAGTATTGAACAGGA
>Gm05_3 8252318	CACAGAGGCATTGTTTGGAAGACCTGAAATTGTTGTAACACCTGAGGGGGCAGTCTCTAAG GATGAAAACATCAAAATTAGCTTTGGTGGTTTAAAGAAGCTTGTTTTAGAGGCTGTGACTTT TGGTTCTTTTCTCTGGGATGTTGAGAGCTACGTGGACTCAAGGTACCATTTTGTGT[T/G]TAA CTGAGTAAAGCTTTGTATGTCTATCTTCTTTAGAGACCACAGGTGGGAGCCTTGTGCATTCA



	GTTGTCCTTTTGATTCTGTTTGCTCCCATACATCACAAGATACAGTTCATACTTCATAGTGT GTTGGTACTGACTGATGAGTTTAAAGTCCAAATATTGTTTCACTAATTCCTA
>Gm05_3 8252356	CACCTGAGGGGGCAGTCTCTAAGGATGAAAACATCAAAATTAGCTTTGGTGGTTTAAAGAA GCTTGTTTTAGAGGCTGTGACTTTTTGTTCTTTTCTCTGGGATGTTGAGAGCTACGTGGACT CAAGGTACCATTGTTGTTAACTGAGTAAAGCTTTGTATGTCTATCTTCTTTAGA[A/C]ACCA CAGGTGGGAGCCTTGTGCATTAGTTGTCCTTTTGATTCTGTTTGCTCCCATACATCACA GATACAGTTCATACTTCATAGTGTGTTGGTACTGACTGATGAGTTTAAAGTCCAAATATTGTT TCACTAATTCCTATGTATATACTATAGTAATAGTGGCTCTGGATTTTCATCA
>Gm05_3 8278658	AGGTTCCCTCCCTTCAGATCTGGCCTCCTGCGTCAACCTCCGCAACCTCTACATTCAGCGC AACTTGCTCACC GGCCAAATTCACCCTTTTTGTTTCATTTGCCGGACCTCGTTCGCTTGAA CATGGGCTTCAACAACCTTCTCCGGCCCCCTTCCGTCCGCTTCAACAACCTTACCC[G/T]TT TGAAAACCTTTGTTTCTCGAAAACAACCAGCTCTCCGGCCCAATCCCCGACTTGAACAACTC ACCTCGACCAAGTCAACGTCTCCGACAACCTCCTCAACGGCTCTGTCCCTCTTAAGCTTC AGACATTCCTCAGGACTCTTTTCTAGTGAACCTCCCTCTGTGGCCGACCACTCTC
>Gm05_3 8279366	GGAAGGGAACGCGAAGAAGTTGGTGTGTTTTTGGGAATGCGGCGAGGCGTTGATTTGGA AGATTTGCTCAGGGCTTCGGCGGAGGTTTTGGGAAAGGGACTTTCCGGACGGCGTACAA GGCGGTGTTGGAGGCGGGGCCGGTGGTGCCGTGAAGAGGTTGAAGGATGTGACAATTT[ A/G]CGAGAAGGAATTTAAGGAGAAGATTGAGGCGGTGGGAGCGATGGATCACGAGAGTTT GGTGCCTCTCAGGGCTTACTATTTACGACGGGATGAGAAGCTCCTGTCTATGATTATATG CCCATGGGAAGCTTGTCTGCCCTTTTACATGGTCCGTCTTCTTCTCTTTTTTGTGCTTGCT
>Gm05_3 8280124	CACGAGGGCCTAATGTTTCTCATGGAAATATCAAGTCATCAAACATCCTCTTAACCAAATCA TATGATGCCAGAGTGTCTGACTTTGGCCTTGACACACCTTGTTGGCCCCCTCCTTACCCCTA ACCGTGTGGCCGGCTACCGTGCGCCGGAGGTGACTGATCCTCGCAAGGTGTCTCAG[G/A] TGGCAGATGTGTACAGCTTTGGTGTGTTGCTCTTGGAACTTCTGACTGGGAAGGCACCTAC CCATGCTCTCCTGAATGAGGAAGGAGTGGACCTCCCCAGATGGGTTCAATCCGTGGTTAG AGAAGAGTGGACTTCTGAGGTCTTTGATCTTGAGCTCCTTAGGTATCAAATGTAGAAG
>Gm05_3 8280387	GGAGTGGACCTCCCCAGATGGGTTCAATCCGTGGTTAGAGAAGAGTGGACTTCTGAGGTC TTTGATCTTGAGCTCCTTAGGTATCAAATGTAGAAGAGGAGATGGTTCAGTTGTTGCAACT TGCAGTTGATTGTGCAGCACAATACCCTGACAAGCGCCCTTCAATGTCTGAAGTGGT[G/A]A GGAGCATAACAAGAGTTGCGAAGGTCTAGCTTGAAAGAGGACCAGGACCAAATCCAACATG ACAATGATATACTATTATAGAAGACATGTATCTTCTCCCTCGACTTTGCTTTGTGAGAGTCG GAGCCCATATTTTACTTGCTAAATCCTGTGCGTATGTTGCTTGCAATTCTGTTGGCT
>Gm05_3 8329040	AAATCCGAGGAGGACCTCGACCCCGGCTCATGGCGCCCCATCTTCGAGCCTCCCGCCGGA GATCCCCAACCCCTCCCCGAATCCGACGCCGCTACTACTCCGCCGTCCACAAGCTCATG TCCGGCGAGCCCGCCCTGATCCAGGACGCCGGCGAGGAGATCGCCGGCCTGGCCGTGAG [T/C]GGGCACCCTGCGGCGCAATCGGTGCTAGGGTTTTTGTGGAAATGGGGCTTTTTAG GAGCGGAGCAAGGGGAAGGCCTTTCTGTACCACCACTTCGCCGCGGAGGGAGGCAACAT GCAATCGAAGATGGCATTGGCCTACAGCTACACGAGGCAGGACATGTTGGAGAAAGGTGT AAAG
>Gm05_3 8386822	GGAAGCCCACAGGCCTGCAAGCCCACCGTCTATTCCAAGATCTTCAAGACCGCATGCCCC AAGGCCTACTCCTATGCCTATGATGACCCACCCAGCATTGCTACTTGACCAAAGCTAACT ATTTCTCACCCTTCTGCCCCCATCGCCCTGATCCATAATTCATCTTGATGCTTGTT[C/T]T ACTTTATTAGACCATTTGATCTATCATCTATGTATCCCTAAATGGTATTATTATTAATGGT GCTTTGATCAATATTACTACCTTTTTTCGTGAGGTGCGCTAATGTCCTCAGTTTATGCTATA GTCGATGGCCAACCACACTCTATTTCTATCCCCACGGTCTAATTGATGTGTA
>Gm05_3 8402327	CATAGGAAAGTACATAGAACAAGTCTACAAGGACCTTCCACCCACTCACCTGCTTTGTTG ACTCACCATCTCAACCGATTGAAATCCTCTGCCCTCCTCGTCTCCTCGTCAAGAAATCCTACAA GCTTCTGGATCTCATCATCTGCCTGCGCTCCAGCCCCAAAAGACCCGTGGTCGTC[G/A]T CCCAAGCCCCAAGCCCCACCCGCTGTGCCTGCTGCAGCCCTCACCGACAGCCCTGCTCTT CTATCCGCCAAAAGAGGCCCGGTGCTCCTAAAAAGATTGGTGTGAGCCCGCCAGGCCCA AGGGGTCGTCCACCCGGGACTGGGAGATCTAAGCTGCCCAAGAGGCCTGGCCGTCCCC
>Gm05_3 8431633	ATGCTCTGTGATATATATCATCATGCTCTACTTGATGAGTTCTCAACAGCCCGTGATCTGCT TCTTATGAGCCATTTGCAAGATAGTGTTCAGCATATGGACATTTCAACGCAGATACTTTTTAA CAGGGCCATGTCACAGTTAGGTTTATGTGCTTTTTCGGGTTGGACTGATTTCTGA[G/A]GCAC ATGGCTGTTTGTCTGAACTCTACTCTGGTGGAAAGGTGAAGGAGTTGCTGGCACAAGGCG TGTCACAAAGTCGTTATCACGAGAAGACTCCAGAACAGGTAAGTATCCCCCTCCCCCTCT TCATTTTCCGAATGATTGATATGGTGAAGTTTATTAGGTGCTGGTCTTAGACTC

>Gm05_3 8432647	CATCAATCTAGAGCTCCTGGAGGCAGTACATCTTGTCTGCTATGCTGCTGGAAGTTCCC AATATGGCAGCCAATGTTTCATGATGCCAAACGGAAGGTCATTAGTAAGACTTTCCGTCGCC TGCTTGAGGTAAGTGAGAAACAAACATTTACTGGTCCACCGGAAAATGTCAGAGATC[A/T]T GTTATGGCTGCCACAAGGGTTCTTAACAAGGGAGACTTCCAGAAGGCTTTTGACATTATTG TGTCTCTTGATGTGTGGAATTTGTGAGAAATCGAGACACCGTACTTGAAATGCTGAAGGA CAAAATCAAGGAGGAGGCATTGAGGACATACCTCTTTACCTTCTCTTCATCTTATGA
>Gm05_3 8432746	CATTAGTAAGACTTTCCGTCGCTGCTTGAGGTAAGTGAGAAACAAACATTTACTGGTCCAC CGGAAAATGTCAGAGATCATGTTATGGCTGCCACAAGGGTTCTTAACAAGGGAGACTTCCA GAAGGCTTTTGACATTATTGTGTCTCTTGATGTGTGGAATTTGTGAGAAATCGAG[A/G]CAC CGTACTTGAAATGCTGAAGGACAAAATCAAGGAGGAGGCATTGAGGACATACCTCTTTACC TTCTCTTCATCTTATGAATCGTTGAGCCTGGATCAACTCACAAAATTTTTGATCTCTCTGTT TGTCGCACTCATAGCATTGTGAGTAGGATGATGATCAATGAGGAGCTTCATGC
>Gm05_3 8433152	CACTCCAGGCTTCAGGCATTGGCATTCCAATTGACCGAGAAATATCTGTCTTGCAGAGA GTAATGAAAAAGCAGCAGAGGCCAGGTTAGGTGGTGGTGGATTGGATTCGCTCTAAGGC GTAGAGATGGCCAGGACTATGCTGCCGCCGCGGCTGCCGGCAGTGGTACAGCCTCTTC[C/ A]GGTGGTAGGTGGCAAGACCTGTCCCTTTCTCAGCCAAGGCAAGGCAGTGGGCGTGACG GGTATGGTGGAAGGCCAATGGCATTGGGCCAAGCAGCTGGGAGTGGCTATTCAAGGGGAC GAGGAAGGGGTTCTGATGGAGGTTCAAGGAGGACTGCTCAAAGGGGTTCAAGCATTGAGGG GA
>Gm05_3 8435563	TGTCAATTCTTCCCAATTTTTTCATGCTCAATGTATATTGCCTTTAACATTGGGATGCAGCTTA GTGTCAAGACTATTAGTTAACAAAGTGCTTTCTGAATGTTTTATAGTTGATCTAATTTCTGA ATATGATTATCAGAAATTTGCAGTGGGGAGAGGACCGACGCTTTGATGAAATG[T/C]GCAGT AATTTGGGTAGACTGGCCATATTTTGGATATTTTCAGGTTTCTTAGATTTTAATCATTTTTTTA AGATTTAGCATGCTGATCTTGGTTTTTTCTGAGATTTATTCTCTTTTGATGCTTGTAGTAGAG GATTTACTTCAATTTTTTGCAGGCTGCTGGGTGTGGGCTGTGAGTTT
>Gm05_3 8437624	TTAGGATGGTCAAATTTTGGTGTGTGACTATTACAATATCGAATTGTTTAGGATGGTCAAATT GTTTAGGATGGTCAAATTTTGTGTCTCTGGAAATGCTTTAAAACTCCTTATTGCATTCCTTA ATTGTAGGTGAAATGGTATTAAATTTTTTAGTATCTGCTCAAGGTGCAGTGCA[G/A]ACAGTG GGAAAATCAGTGATGCTCTGAAAATGGCTAGCACTTGGTTCTTTGAATTTGTTTGGTTCTC GTCTTACGTCAGACTGAAGTTGCAGCTGGTATCATCATCCGCCATGGTTAGGGATGTGGCT TTGTAATGCTATTGCAGCTTTCCTTCTAGATTTAATCTTTCTAATTTGCTT
>Gm05_3 8437721	TGCTTTAAAACTCCTTATTGCATTCCTTAATTGTAGGTGAAATGGTATTAAATTTTTTAGTATC TGCTCAAGGTGCAGTGCAAACAGTGGGAAAATCAGTGATGCTCTGAAAATGGCTAGCACT TGGTTCTTTGAATTTGTTTGGTTCTGTCTTACGTCAGACTGAAGTTGCAGCTG[T/G]TATCA TCATCCGCCATGGTTAGGGATGTGGCTTTGTAATGCTATTGCAGCTTTCCTTCTAGATTTAA TCTTTCTAATTTGCTTGTCTCTCTTAGATATGTATAAATTTACGTACTATATGTTGTGTGC AAGTACTATGGCTTGTGCAGGTAACATAATGGATCAATCATGTTTTTCAT
>Gm05_3 8437894	CAGCTGGTATCATCATCCGCCATGGTTAGGGATGTGGCTTTGTAATGCTATTGCAGCTTTC CTTCTAGATTTAATCTTTCTAATTTGCTTGTCTCTCTTAGATATGTATAAATTTACGTACTAT ATATGTTGTGTGCAAGTACTATGGCTTGTGCAGGTAACATAATGGATCAATCATG[C/T]TTTT ATCTTATTTCTTCTATTGCAACTCTCATTGTGGACCTGCAGCATGCCAACATCGGCAGATAT TGAACACAAATGTTTTGTTGATGTATCCTAATATTGTATTTAGTTAAATTCATTTCTGAATACA ATGTAGTTATAAACTAACGGAATGCTTCTGTCTGGTGAATGTGCTGAGA
>Gm05_3 8437928	TGGCTTTGTAATGCTATTGCAGCTTTCCTTCTAGATTTAATCTTTCTAATTTGCTTGTCTCTC TTAGATATGTATAAATTTACGTACTATATATGTTGTGTGCAAGTACTATGGCTTGTGCAGGTA ACTAATGGATCAATCATGTTTTTCATCTTATTTCTTCTATTGCAACTCTCAT[A/G]GTGGACCT GCAGCATGCCAACATCGGCAGATATTGAACACAAATGTTTTGTTGATGTATCCTAATATTGT ATTTAGTTAAATTCATTTCTGAATACAATGTAGTTATAAACTAACGGAATGCTTCTGTCTGGT GAATGTGCTGAGAAAGTCATCTGTCTTGGTCTGCAAGGGTGTGTT
>Gm05_3 8588041	CCCCACCGTCGTCTCCGACGCCCTCCACAAGGCTGCCGTGAAGGCCGTCGACGTTCTCAC CGCCATGGCCGTCCCGTCGAGCTCTCTGACCGGACTCCCTCGTGAAGTCCGCTAGCAC ATCCCTCAACAGCAAGGTCGTCAGCCAGTACTCCACGCTCCTGGCTCCCCTCGCCGTCG[C/ T]CGCCGTTCTTTCCGTCGTCGATGCCGCCAAGCCCGACATGGTTCGACCTCCGCGATGTG AAGATCGTGAAGAAGCTCGGCGGCACCGTCGACGACACCGAGCTCGTGAAGGTCTCGTC TTTGACAAGAAGGTCAGCCATGCCGCCGGGGGACCCACCCGCATGGAGAACGCCAAGATC GC
>Gm05_3	TGCTGAGGCGCTTGAAGTTATCCCTATACTCTGGCTGAGAATGCTGGTTTGAACCCGATT

8589019	GCCATTGTTACCGAGCTGAGGAATCGTCATGCACAGGGTGAGATAAATGCTGGAATAAATG TGAGGAAGGGTCAGATTACCAACATCTTGGAGGAGAATGTGGTGCAGCCCTTCTTG[G/A] TAGCACAAAGTGCATCATGTTGGCGACAGAGTGTGTGCGGATGATTTTGAAGATTGATGAT ATTGTAAGTGTGAGGTAGAGTTGATTGAACAATGTGGAATTTAGAGAGAACTGAATCACCC ATTATCTTCTTTTTTTTTTTTTTTGTACTTCGTGCTTTAATTTTATGTTGGTAGTTT
>Gm05_3 8596037	GTAGATAAAATAGAGGTAGAGGTGTGGTAGATTTGAGAGCAACATTGAAATGCATGCCAAG ACAGACTCGGAGGTAACAAGCCTGGATGCGTCGTCCAGCACAAGGTCTCCTCGGCGAGCA GTGTACTACGTTTCAGAGCCCTTCCCACGATGGGGAGAAAACGACGACGTCGTTGCACT[G/ C]CACCCTGTTCTCAGCCCCATGGGTTCCCTCCTCACTCTCACTCCTCCTCCAGCCGCTT CTCCGCTTCACGCCACCGCAATAACCATAATAATAAATCTTGAAGGGCATCGACGTCATC GAAGAAGAGGGTCTTCTCCAATCCGAATTAGATCGCCAACATTCTCTCTCTCGTCGATA
>Gm05_3 8696065	AATACCACAGTACAGTGC GTTCAGGAAAAATGAAAAACCAAGCCCCAGAAGTTCGATTTCAA GTGGGTAACAGAGCAGTAGAGTTGATCCATTACACGTAAGGTAGCTACTCTCCTCAGAAGA ACCAATTTTTCATAAGCACCATGACGAATATCGAATATTTGAACCTAGTGGAAATCAGG[G/T]TC AAACTCATGAGGTGTATTCAAATGGTTGAGGAGGTTGAGGTTCTCATCCACAGCCATGGC TGATGAAGAACTCTGCAATCCTGTTGCTGACCCACCAGCAGCAGAACTGGTAGATGAGGG CGTGTCTAGTTAGAGCAAGCACTTCTGGCTCAGTAGGATGCAAGTCTCTGAGAAGCAC
>Gm05_3 8765188	GGAGTTTGACAATGATAACAAATTTCCCACTAAACATAAATCTGCAGATCTGAGCATCCCTAA ACCCACTGAAAAGGCGATTTTAAATTTCTTTCTTGGTTGAGTATGGCAGGTATTGCAATGG AACATTTTCTCTGGACTAAAGAAAACCAAGCATGTGCGACCGAGAGGAATAGCAC[G/A]CTA GGTCTTCTAACTGTTTACCCAGTAAAGTCGGGAAGTTTTCCACTGTTGATTTTCTCACC GG GAAATTCAAGAATGGAAGAAAGAATGCTACTCTGTGTATATCTCTGTATATTAAGTGTAAT TTGTTTATACTTGTTAAAGGGTGCTTTATCAAATCTTGTCAACCGGTAAC TTT
>Gm05_3 8863867	AGATTCCTCGTCCCTGGGAGCGTCGTTCTCCTGCGGTA CTGAGCCACACGAGGGCCAC GTGGACGAGGCACTGCAGCGCGTAGCCGAGATCCACCAGCGGATCGGCGTGTGGGGT TCTCCTTGACGGTGCAGGCGAGCATCACCGCCGAGACCAGCACGAATGCCATGTTCCACG[ A/T]TATGTCGAGCGCCACCACCGGCTTCGAGTAGGTCCAGTCGGCGCGGCGCTCCTCCAG CTCACGCGCGCGCTCTCCCGGACCACGGCGGAGTGGCCGCGGCGGCGGAGGAGGAGC GCTAGGATGGGGAGGCGGGCGGCGGCATCGGGGCGCGGAGAAGGAGCGGCGCGTGG GACTGAGG
>Gm05_3 8896514	ATCTCTTGACAGATTGTTCTCTGCAGCGATGTATAGCGCAGTCTCTCCAGCATCGTTCTGAAG AGACATCACATCAGAGAGCGATGACCATTCTTCATTCTTCACTTTCTCCACCAGTTTCTTCA CCCCATCAAAGTCCCCAGATCCAACGCTCGAAAATATTGATTGGTACGCTAAGAA[G/A]CGC ACACCCTTTGCCTCCATCATTCACTTTCTCTCTCTCAAGTTTCAATTCAACAAACAAAA AACAAAGACAGGGGTTGGTTGTTGTTTTCACTGTTTCAGCTTCTCCTCCATCATGATTGCA GAAAAGGGTGTCCTAAAGTATGAACTTTTTATTTTTCTTTGTATCAAAAA
>Gm05_3 8926542	CATTTAACTCTTGTACTTGTTATGTTATGTTGAGCATTTGTCTATGGAATTTCTTGACCTTT ACCTTTGTATTAGTTTTTTATGCCTACATAATTTGTATGTAATTGGTTGGGCAGGTTTACTT GGACCCAAAGGAGCGAAACAACACTGAGTACAAGTTGGAGACTTTTGTCTGCAG[C/T]ATACA GGAAGTTATCTGGAAGATGTTGTGTTTGAAGTATCCTGTCTCAGAGTCTTAGTTTCACTTT TGTTAGCTTTGAGAATTTGTTGCGAGAATTGCATATGGAGTATGCTGGTTAAATTTTGGTG ACAAATTGACTTTCAATCTGGAATTCGTTTTCTGATTTTTGGAAGTACTAT
>Gm05_3 9038383	GATAGTTTAGGCTATACAGTCATCTAGACAGCAACTACATCCAATTGTGATTCATATTATCC ATTTCTATAAATTTGTTTGGCTCAAAAATGTGATTAGCATAAGTAGGTTTACACATTGAATTA ATTAACCTCCTCTTCTTGATATACATAAATTATCCTTAGAACAATACAAAA[T/C]TGGGTG TGGAGTGTGAGTATGTATCTATGTAATTGACAGATCTTTCAGCTGCTGGGGATCCACC TCAGATGGTGATCTCGTGAGGGCACACTGTGCAGTGGTTGTTTTGAAAAGCAATGACAT CTCTGATGGAATTAGCCCCGCCAACAGCATAACCAATCTGTCCAACCCAT
>Gm05_3 9082457	TACATAGAATTACTTAATATTTGATTTTTTCATGAGAGTGATTATTATTTGCTGCAGCATGTG GTGCTAGTGGGGCTCTGTATACGCTGATTTGCTGTGTCATTGGCTGTGGCTGCCTATACTC TTGCTTCTACCGCCCCAAGATGAGACGACAGTATGGTCTAAAGGGAAATGGTTGT[G/T]CGG ATTGCTTGATTCATTGCTTCTGCGAGCCCTGCGCCCTCTGTCAAGAATATCGTGAGCTTCAA CACCGTGGATTTGACATGATTATTGGTATGCACTCATTATATATATTAACACTACTCTATT TGCTAAATACCCAGCTATGCAAGTTGGAAAACAAATTTCTTATTTTATAGA
>Gm05_3 9082469	CTTAATATTTGATTTTTTCATGAGAGTGATTATTATTTGCTGCAGCATGTGGTGCTAGTGGG GCTCTGTATACGCTGATTTGCTGTGTCATTGGCTGTGGCTGCCTATACTCTTGCTTCTACCG CCCCAAGATGAGACGACAGTATGGTCTAAAGGGAAATGGTTGTTCCGATTGCTTG[T/G]TTC

	ATTGCTTCTGCGAGCCCTGCGCCCTCTGTCAAGAATATCGTGAGCTTCAACACCGTGGATT TGACATGATTATTGGTATGCACTCATTATATATATATTAACACTACTCTATTGCTAAATACC CAGCTATGCAAAGTTGGAAAACAAATTTCTTATTTTATAGAGAAACTCCTTTT
>Gm05_3 9082823	CCTTTTTTGTAAATTAATTTATTTTTCTTTGAACTTGGTACGGTTAGGGTGGCATGGAAATG TCGAGCAACGAAGTCGGGGAGTAGCCATGACTGCTACAACAGCTCCATCAGTCGAAAATG GCATGAGCCGTTGAATAGTATGCAGTTCTGTAAGACAATGTGTCTACGATGC[C/A]CG TGTTAAACATGAAGAATAATAATGTTTGAAGAATGGTTCATTCTTTTTTTCGATCCCCTTTT CATATTAATAGTTGTCATTTAAATTAATGTTAGACTACTTTTATAATAGATGCGGTATTACAAC GTATCTTTTTCTGTATCCAATTGTGGCAACGAGGATACAATATAAGTTCTT
>Gm05_3 9083038	TGGTTCATTCTTTTTTTCGATCCCCTTTTCATATTAATAGTTGTCATTTAAATTAATGTTAGA CTACTTTTATAATAGATGCGGTATTACAACGTATCTTTTTCTGTATCCAATTGTGGCAACGAG GATACAATATAAGTTCTTATGCATTTATCTTAACCTCCGCCGCTAGGCTGAT[T/A]ATAGTGG TTTACAATGCATCTATTTAAAAATAAAAAAGACAATGGATCTATTTATCCTTTCATCCTG TGTGAATATGTTTACAAAATCTAAGGTTTTGTGAAACATGGAATGACTTTCTTGAAGGTGT GGTTGAAATGTACACTTGACTTATAGAACCAGTAACTCTTGATGTA
>Gm05_3 9085416	GATTTCCGCAATTTCTTTGTAATTGGGGTGAGGTGGTGGTGTGGCTGGGGCTGGGGCTG GGTATACTCTTGCTTCTACCGCTCCTATATGAGACAACAGTACGATTTGCGGGGAAATGCC TGTACGGATTGCTTGATTCAATTTCTTCTGCGAGCCCTGTGCCCTCTGTCAAGAATATC[C/G]T GAGCTTCAATTCGGTGGATTTCACATGACTATTGGTATCTACCATTGCTTAATTCATTATTA TGGCCCCTAATTTGTTTTGCTTTCCTTTTTTCCACATCAAAACAACGCTTCCATTACTCTC ACTTTCATTTTTCTGTTTTGTTACTCCTTTCTTCAATTGATACAGCTCATAAA
>Gm05_4 0859886	AGAGTTGGATTTAAATCTAGTTTTATTGTTATGGTCTCCGTAGTTAACAAATTTTATCAGCA TGTTACATTATTACTCCATTCTACGTTAGGTTGTTGACGAATTGAGTTATTCTGTCATCAGTT GAAATTGAGTTGAGAGAATATCCAAATCGTTTGGCAATCTGGTGGTGGAGGGG[G/A]TATCA AGGTAGGAAAATTGTACTTTGATTAGGTAGTTGGTGATGTTCTTTTACCATTCTTCCCTCCCTC TGTAATATTACATCTGATGAAACGTTTAGCAGTTTGTGAATACAGGTTTGTACCTGTAGT GTAAGGTTGTCGTCAGTGTGCCATGTATTCTATTTAATAATAAAACTTCA
>Gm06_5 092543	TGGAAATGCAGAAAAAGTATGTCTAAATTAAGATTGTGTTCCAGGCTGAATGTTAAATGCAT GAGATAAAACAATACAAATAAGGCATTTAGGGGAAAAAATATATTAGTCTCACAGGTATTG TAAAGCAGGTTCTGCATCTGGCTTCTGAAGTCGCTCAAGAAGTGCATAGGCAGCC[T/C]ATG GGTGTCTGACTGCTCAACCAATTTGGGCACCAGAGCAACTAGGTCTGCTTGTAATGCTT AGGAGCCTTGTCGAATACAAGAGAAATTTTTGTGCAAGCTGAGGTGCGCGTCTTGGTTCT GGACAGCCTTGTTGAACAGCCCCGTCCAAGCTCTTAGTGAGTTTACAATCAAAAC
>Gm06_1 1230990	ACATATATTATTATTCAAAAAGGATTTGTGCTATGCCGTTGTCTATTGATTGTCAGTGTTC TGATCTTAAGGCTTTACCTGAAGAAAAATGCAATGTACATCTTTTTATTGTGTGCGTCTTCTA TTGCAAAACCATATCTGTTGATTTGTTTTCTTGACTAAATGGACTTTAAGAT[T/C]CAGGTG GTGCTGAGGGTGGGAAGCAGAGCAGAAGTGAGAAGAAGAGCCGAAAAGCAATGTTGAAGT TGGGACTGAAACCTGTTACTGGCGTTAGTAGGGTCACAATCAAGAGGACAAAGAATGTGAG CATTGTTATTGTTCTATCTGTGATCTATCTGGGAGGCGGATGGATTTTAACCA
>Gm06_1 6005493	TCGACCAAGGATTACAGCACCACTAATAATGGTGCCACAGCTAAGACAAGATTTTGGCAA CCCTCTTTACAAAGTATCTCAGAGGCTGCCAGCTGATCTTTGTTTTCAATGTTACCCACATG TGGTGTGAGGGACAGAGAAGCAACTGAGGAAAAAATAAACAGCAAAAACAAAA[G/C]G AAAAAAAACCATGATTGCTTGGTCATGTGAAAATTAATATAGTACACTGAGGTTCCAGGGG AGGCCAGAAACACAGAGATGATAAAATCTCCTCTGAGTTGAGGCATAATTGTGTATAGTA AAGTTGACATGTCTATTTCTTTTAGTGTAATTAGATATTACTCTCTAATGAAAT
>Gm07_1 262867	AAAAAAAAAAGTACGTTGACATCAGTCAACATTGCAGGGTAGGTGAGTAGAAATTTTTGA CAAGGATTTTTATTTTGTGGGTAATAACGTGGAATGTTTAGGGAACATATCTTACTAACC CTGTCTATTTTCTACGCGAAACAGAAATTTTTGAAAGAAGAAATGACTTACTT[T/C]TAGC GTGTAGTCGGTAATGAGACTGCTTCTGTCCCTTCCATTGCAGTTGCCTAAAATGCAAGAAT TCCATTGATTAATAAATTGGCATAATTATTTACTATTACTCTATTAGTTAGTCCATTAAGTGTT ATTTACATAATAGTTATTACTTAAATGGGAAATTAATGAAAATATACCTT
>Gm07_1 262870	AAAAAAAAAAGTACGTTGACATCAGTCAACATTGCAGGGTAGGTGAGTAGAAATTTTTGACAA GGATTTTTATTTTGTGGGTAATAACGTGGAATGTTTAGGGAACATATCTTACTAACCCTG TCTCTATTTTCTACGCGAAACAGAAATTTTTGAAAGAAGAAATGACTTACTTTA[T/C]CGTGT AGTCGGTAATGAGACTGCTTCTGTCCCTTCCATTGCAGTTGCCTAAAATGCAAGAATTCCA TTGATTAATAAATTGGCATAATTATTTACTATTACTCTATTAGTTAGTCCATTAAGTGTTATT ACATAATAGTTATTACTTAAATGGGAAATTAATGAAAATATACCTTCAT

>Gm07_4 060462	AATTCATCAGAAGCTTTTAATTTAACTACTGATGAAGAGAGAACCATGGCAAATTTTGTTTT ATTATACATTAAGGCCTCTTGATAATTTGGATGAGTTTAAAATGGTAAAAATTTACCTGCAT GAGAACCAAGACATCAATTGGATCATTGTCTTCACACAGAGTTCTTGGGATGA[G/C]ACCAT AGTTATGAGGATAAACTGATGAGTACAAAACCCGATCAACCTGACCATTAATTAGAAGG TAAGAAGAATATAATGATAGACGAGTAATGGCATCCACAAGTCAGAATTGCGAAGGGTCAA GATTTGAGAATCCACAAGAATTGTAGAAAAAACATGATCACAAAATATCATC
>Gm07_4 868495	TCATATTATCTGCTCCCTACACAATAGGGTGGGTAGGAATTGGGTTTTCTAGGGATGGTATG ATGGTTGGTTCCAGTGCAATGGTGGGATGGATTAGTAAACATGGTCATGCAAAAATCAAGC AATTTTACCTAAGGGGTAGGAAGCAATCAGAAGTCATAATAGACAAAGGTGAACTC[T/G]TC CTCAATAATATCCCTGCGGCTGTGGCGACTAATGGAGCTGAAATTCATATAGCATTTTCAGTT GCAGATGACAACCCCATTTGAGAAACAACCAATTCTGTTGGCTTTTGGTAGCAAATATCCAC AGAACCACCACCTCTCCAAGCATGAAGACAAAACAGCTATCGTGTGTTGATTTT
>Gm07_5 666664	AAATATTTAACTCGACATACGGTGTGAATATCCTTTCTCCACCTTATTCTTTTTATGGTATCC TTTTCCACCTTATAAATAGTTCATTATTTCAATTATATAAAAAGAGCTGGATCTATAACTGCC CTAACCTGCTTTTTCTTCATTGTTGGTTTTCTCTTCGTAGCTTACACCCAACAT[A/C]GAGCAC CTGGCACTCGGTGTGCATGAAGTGAAGATGATTTTGGTGGAGAATTCCAGGGAAACCACT TAAACAAGTTAAAAGTTCTTGCTCTGTTCTTTCATACTGAGTCCGATGTGTTTCTACAACGG GTGCCCAATATAGAGAAGCTTGAGGTGCTTGATGGTTTCTTCACAGAGACT
>Gm07_7 162763	TCAACTCCCCAAAGCATTTTCGTCGCTTACTACGCCCTTCCTCATCTCTGGGTGCCTAGGTAT CCACCATAAGCCTTTCTTATATAGCTCTCCACAACCCCGTTTTACGGTTTAGGCTGCTCC CATTTGCTCGCCGCTACTACGGGAATCACTTTTGCTTTCTTTTCTCTAGCTAC[G/T]AAGA TGTTTCAGTTCACCGGGTGTCTCTTGCTGCCCCGTGGATTACAGCACCAGTTCGAAAGGTT GACCTATATCGGGAATCTCCGGATCTACGCTTCTGGGGTATCTAATCTCATTGCTCCCTA GCTTCGTCTCTCGGTGTCAGTGTGCGGCCAGCAAAGTGCTTTCGCCGTTGGT
>Gm07_1 4560429	TGGATTAATAAAAAACATTTTCAAGTGTATTATTTTACCAATAACTATTATAAAGTATCACAA CGTATATATAAGTATTGATTGGACTATATATGAAGGTAAAAACTAGCACATAATAATAGCA TTTTAATTTAAATAGATGTAATGAAAGAATGAGTAGTGGTAGTAGAGGTT[A/G]CCGGAAC GGAAGGGAAATTGAAGAAGAGATTGGCGTCGTCGTGGGAGATTAATGGAGAGTTTTGCGG AGGAGGAAGGTGGTGGTGGGTGTCGGTGGTGATGTTGGAGAAATCTAGAAAATCATCAAT ATGTAAGCAATCTGAAGGGGTGGAAGAAGGAACCGTACAAGTCCATTGTTTTTC
>Gm07_3 8744579	TTGTTCTATATTTACAATGTGATGCTCAATCGACAATGGATTTAACTTAAAAATATCAGTAGA TCAGGATTGTAGAGAGAATAGTATAGTAGTAAAAAAGAAACACATCAGGATTCATCTACAAC TCCTCCTCAATTCATCTCCTATTCCTCCCCCCCCCCCCCCCCCCCCCGCAG[G/A]TTCC TAAGACCTGCACTTGAGAGTAGTTTTTGGGTGACTCTCAATCGCCAGTGGAATCTCCCTC TGCTACTTCGTGTCTGTGATTACCTTCACTATTAGTGTGTCGTTATCGTATTTCCCCCATG TCTCTTGATGAGACCTTTCTTCAACATTGCCATTATCGCCACCTACTCTC
>Gm08_5 85476	GCCTTTCCAGGGCGAGGCTGTAAACAATGGTGTGCTGATCCAACATGATGGTCCATCATATG TGGTGTCTCAGCATATGAGGCGGTGTTCTAGCAAAGCTTGACATCTGTGGAAGGCCCGGA GAAGAAGCATCATTAACAAATGATGGTGAATTTCTGCCAAAGCATGGTACTTGATGAA[T/C]GT TGCTGTTGCAAATTGGAGCTGTTCCACATATGATAAAGTCCCTGATGTGGGCGTATCCCAT TCCTGATGAGCAAAAGTAAACAAAGATACCACATCAAGATTAGATGTAAACTGTTAACT CACATTACAAATTGGTTCAGCTTACAATTACCCAGATCCAACCACTTGAACCTGA
>Gm08_5 86892	AGCCCATTCCCTGAATATGCCTGCCATCAGAGGCTTCTGTTGCTCCAGCACCTATATTGTTA GCCATGTTGCTAATGGAGCTAGAAAAGTTATAAGGACTACCATTAGCTAACTATCACGATA CTCAGGTAAGGAATGAGGATGAAATCTCGAACCAGATGCAAAATTCATTGCATCA[T/C]CAA AGTTACTTGACTCACAAACACTGGACACTTTTCCGGCAGGTGCCCTCTCACAGTGTGTG AATGCCAGAATTCACATGAATGAATGCATTATCCATAAATGCATTCAAAGGTTGCCTAACTG TAGATTGAAATCTCTGATTATATCCATTTTCTAAGCAACCAGATCCAATTACTC
>Gm08_1 908246	CTCAGAGCAAGAGTGAGAATGTGAGTGAGGCTTCGGGAGAGAGAGATTCTGTGTATTTGAA AGACTTGTTGAAGGAGTATAGAGGGAACTTTACGTGCCTGAGCAACTTTTTGGCACAGAG TTGTCGGAGGAAGAAGAGTTTAACAGAAATGTAACGAGTTGCCAAAATGAGCATT[G/C]G GGAGTTCAGGAAGGCTTTGAGCAAAGACAAAATCAAGTTGATAACTCAAAGGGAGGTGGA GGTCTTTATAGGGATTTTGTGTTGGAGTTGAAGAAAATTCCTGGTGATAAAGCTTGCACAC AACCAATGGTAACAGAAAATGTAACCTTTCTTGTTAGTTATTATGCGCTGTGCAAT
>Gm08_2 035440	ATTTGAAAATTTTAGTCTCACATCGCAATATCCAGGGTGTTTGGTTACGAGTTGGAATCATG TAAACGGTCAAAACACAAATTCCTTGCTTCTTAATTTTCATGTGTTGGAAAGTGTGAACCCGTG TTTGCATCACTGCAAATGGGTTCCAGAACACACATTGTCTTCTCTTCCAATAC[T/G]GCGA

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>Gm08_2 670235	CTTCGCGCCGCGGTCTCCGCCGATACTGGGGCGTCGATTGAGTGGCAGCCTTTGGCGG CGGCGGCGGCGGCGATTTCGGTGGCGAGGGAGGGCTCGGAGGTTGTCATGTCGACGAGG ACGCCGCCGGGGCGGAGGGCGGAGAGGGCACCGGAGGAGGGGTGGAGGAGGACGGAG CGGA[A/T]ATCGGAGGGGTAGCCGACGATGGTGAAGACGACGTCGGAGTTGGCGGCGACG GCGTGAGGGGAGGGGGCGAAGTGGGCTCCGAGGTCGAGGAGGGGCTGGGCCTTGGAGG GGGTGCGGTTGAAGACGGTGAGGGTGTAGCCGGCACGGATGAGGTGGGCGCACATGGAC TGGCCCATGAC
>Gm08_2 798583	TGTAATCAAGAGGCGTCGGAAATCTCAAAGAAAAAATGATGAAACAAATCAAAGTAGCAAT CGGCATTAGATCGGTGAGCTACAGTGACATAAAAAATGAAATAGAATAGAACATATGAAAT GATGAGAGAAAGAAATAGAAAACCTGAAACACCGAGATCATAGCCAAAGAGAGAAC[A/G]G CCAAGGGCCCCAACGAAGCATGTATAGACGAAATACCACTGAACCTTGTGCTCGTAAAGAT GAGCCCTTTTCTGGTCTCCAATCTCTACTCCTCCACCAGCCATTCTCTGTTTCAAAGA GAAAAGGGAAAGGTCCACCAACACAGCCTTACAGGTTACACATACAGAACTCCAG
>Gm08_2 864933	ACAAACAAAATGCTCGCTCACAGAAATGCAAAATTTACATACTACGGTAAGCCTTGATGACT GTAACCTGAAGTGTGGGAGAACAACCTAAAACCTTAAACATCTACATCAATGATATAGACGGC TACTACAACAACCTAATATGCTGTCAATATATATTGCCCACTAAAAAGTTT[C/A]GGAC ACCCCAATCTCGTATATAGAATTATACGTAAGAAATACTCAGGTATTATATGCTTAAATGTGG TCTTCCATCTTTCAGGATTAGAAGACAAAAGCATACAGCATTGCACTAGCTAGTAGCGACTT TTGCACTTTTCCATGGCTCTAGGTGCTGCAATAATCACAAATTTAGATGAC
>Gm08_2 983624	CTTGTCTCTCTAAACATTCGCAGTGGCCGCCCAAGAGCTTTGCGCTTCGCCAGGCTAGAGA GCTAATGCTGCTATTATCCTTCCTTGGATGTCTGAGGTCAAATTATATCATAAGTTTGACTAT TTTCTCATTTCAGGCTAGCACCTGTGTCTTGGTACGATTTTGTACACATGGCT[T/C]AATG AAGATGGTGTTCGTCACCTCATATTTTGTATCCACTGGAATTAGTCTGTTCCCGCATAGTT GACAAATAAATACTATGTATAGATGGAATGGATCCTGAATCGTATGCCATCATTTAAACGTG TATAGTTAGTGTTAATGAGTTGAAACGGATCAACATTTAGTTGTTTTTTG
>Gm08_1 0919109	ATATTCTGGAACTTTTTGTTTATTCTTTTTAAAATATCATATCATACTGATGAAATTTGCA TGTTACATACATAACTATAACTACAATTTTTGTGTGAATCAGATCAACAGTAATAGTGTTCA AGAGCTGCCCAAAAATAAAGAACAGCTAAGCAGCTTGTGTAGACAAGAATCT[G/A]ACAGTT GGAGAGTGGGTTTGATGGTAGTGGGGTTGAGGGTGCTCACTGCTGTGGGCAGTGTTGACG CTGCTGGAGGTTGTTGAATCCATGAATCTTCTTTCATTGCCAAGGCTTGCCCTGGAAATATT TGTCTTGTCTGCTTCCTAGTTTCCCGGATCTTTGAGAAATCAAGTGAGTAG
>Gm08_1 4156009	TGGCCGTGGAGTTTCACCGATGGCTTTGGGGAACACGTGGCGGCCGAGGCTTTTTTCTTG GGAGAGGCGTTGGGTTTCGGTGAAGCGGCAGGCGTAGCTGGCGGCCGCCGAGTGGAACCT TCACGGCGTGACCATGCCGTGCGCGTCGAAGAAGTGGTGCCAGCGACGGGCTCGTAG A[T/G]TGGGTTGCGCGCGTTGCGGACGTAGACGCCCTCAATGCATTTGGGGATTTTCCGG TCACCGGGAGGGACTGATGCGCCGGGTGCTCCGGCACGGGGGCGAAGTTGCCAGCGATT TGGACACGCGGGTCGGCGGTTTTGGGGAGTGGGTGTTGCCGCTCGTGGGAGACCAAGGC TGTTTC
>Gm08_1 4997737	CGATTTGTATGCATGCTCAATTTCTTTCTCAGATTATAGTTCTGCAATATATCAATTACGCC CATGTAGAGGACAACATCATATACCTCAAAAAGCTCTACTTCTTTTGCCTCCACCTCCTCCT CAACCTTCCGAGTAGCTTGAGCTGGCATGTTCACTCCAAGCTGCACTCGCAACC[C/T]GAGT CACAAAATCAACGATCCAAAAGATCGGGCATTATACAAAATGGCAAACAGCCTATGGCAT GCATGTCCCTGAATTAAGCTCGCAGCTGTCATAACATATCTTACAAGAAAAACATCTTTTA TGAGAATGAGAGGATTAATCTGGACAGTCATATATGGAGGGTTAAGGGTTTA
>Gm08_3 2448434	CTTAAATTTTTTTGGGGAATTTCCACATTTCTGTTCTTGAGGGGGTTATTGTTTTGAAACGC GTGTTGGTTCCGCATTTTGTGGGGGTTTTGGGGGTTGGGGGGTTCTGGGGGGGGGTGGG GGGCTTTCCGTTTATGGGTGGGGGGGGGAAGGGGGGGGGGGGGGGGGGGGTTAT[ A/G]AAGGGGGGGGTGGGGGGGGGGGGGGGGGGGGTCTGTGGATGTTTCTGTAGACAA GTGGCCTCAGATATCTTAAGAAGGGGGGGGGGGGGTTGAATTAAGATATCCAACTACTTC CCCAAGTAAAAACCTATTTCACTTTTTATTTAAGTTATAAATTTCTTAACAATGAACTTCTT
>Gm08_3 2448435	TTAAATTTTTTTGGGGAATTTCCACATTTCTGTTCTTGAGGGGGTTATTGTTTTGAAACGCG TGTTGGTTCCGCATTTTGTGGGGGTTTTGGGGGTTGGGGGGTTCTGGGGGGGGGTGGGG GGCTTTCCGTTTATGGGTGGGGGGGGGAAGGGGGGGGGGGGGGGGGGGGTTATG[ C/T]AGGGGGGGGTGGGGGGGGGGGGGGGGGGGGGTTCTGTGGATGTTTCTGTTAGACAAG TGGCCTCAGATATCTTAAGAAGGGGGGGGGGGGGTTGAATTAAGATATTCCAACTACTTCC CCAAGTAAAAACCTATTTCACTTTTTATTTAAGTTATAAATTTCTTAACAATGAACTTCTTA

>Gm08_3 2753756	ATCGAGATTTTGCATCAAGGAAGGGTGATCCCATTGCCATTCCCTCCTGGGGTTGGTGGAG GACGCATAGTTGGAGATGCCAACCTTGATACTTTTGAGGTGATTACGGCCGAAAAAGCAAT TGACGAAAAACAATGTTGGTAATCGGATGCTTCGCAACATGGGCTGGCAGGAAGGATTG[G/A ]TACCGTTTTGTTCCCTCCTCATTACATGAGTTTTTACGTATTACATTTTTTTGTTATTCAAGTT GTTTGGTAGAGCTTGATATATTATTCCAAATTTCTCTTTTCAGGGATATGCTAGTACTTTGA TCCTAGACCTAGATAATATTTTCTCTAGTTTAGATATATGTGTATATAATACA
>Gm08_4 2892150	CCTCTGTTGAGATTAAGCTTTCTTTAAATTCATCTGACATATGAAAACAATGGAAAACCTTA AGCTTTGGTAACCTTGTCATACCACGAGGCACCTCTGTCAGTTCGCGAATCCTGAATAACTT GAGCTTTTCGAGAGAAGGCAATGCTCCATCTTCAATTACGATGGATTTCAATTG[G/C]TACAA ATCTGCCAGTAGTATTTGCTTTAGGTTTGGAAACCCCTGTTTGGAACTGCACAACCTTAC TATTGTATGCATGAAGGAGTATACTAAGGTGCGTCAAATTTGGTAGATCTTTAAGTAGTGGC AATGGATCATGTGTCAAGTCGGTGAATGACAAGGACAACGTAACAAGATT
>Gm08_4 3848399	CTGTTTGAGTCTTTGTCATACTTGTACTAAATGTTTTTTTTCAATCTCCAGGTAGGCTTCAG TTGTGCAATTTAACAGAGTGTGCTTAAGAAATAGTAATTTTGATATTTTACTGTAAATAAGAT CAGTATGGGTGAGGAAGAACCAGCGAATGAAGTTTCTAAGATTGACTCAGATG[G/T]GAAGA GCTTGACAGGAGAATACTTTGGATGAAATTACTGAGAAAAAGGACTTGGAACTGATGGAGG AAAAGTCTTGGAGAACAATGGAATCAAAGAGTTGAAAGAAAATATTGTAAAAGAAATTGAAG ATAAGCAAGCTGATAGTATGGAAAAGGTAGAAGAGAGCAAGAAAGTTGATGT
>Gm08_4 3848818	AGAAAGATGGGGTGGGAAGAAGTAAACAAGGATATGAAAGAAGATGGGGTGGGAGAAGTAA AAGAGGATAAGGAAGTTGATGGCGTAGAAGTAAAGGAGGGTAAGGAAGTTGATGGTGTGA AAGAAGTCAAGGAGGATACAGAAAATTGATGGTGTGAAAGAGGTCAAGGAGGATAAGGA[G /A]GTTGGTGGTGTATCTGGTGTGAAAGAAGTCGAGGAGGATAAGGAAGTTGATGGTGTGAA AAAAGTAGAAGATAAAAAAGGTGATGAGTTGAAAGAAATAGAAGAGGATAAGAAAGATGAT CATATCTGAGAATGATAAAATGGATGAAGACACTGAGGTTAAGGAAACAATTGAAGGT
>Gm08_4 3870090	ATTGTAGCTAGTTTTATCGTTTTATTTTCCCCTCTTATTTGCACTTTGCTGTTTGAGTGTG TCATGCCTATACTAAATGATGTGTTTTTTTAAATTTTCCAGGTAGGCTTCAGTTGTGCTATTTA ACAGTGTAGGGCACTGCGTGTTCGAAATAATAAGTTTGTATAATTACTGT[A/T]ATTAATAT CAGTATGGGTGAGGAAGAACCAGCGAATGAAGTTTCCAAGACTGATTCAAATGGAAAGAGC TTGCAGGAGAAAACTTTGGATGAACTACTGAGAAAAAGGACTTGGAACTGATGGAGGAA AAGTCTTGGAGAACAATGGAATCAAAGAGAATATTGTAAAAGAAATTGAA
>Gm08_4 3883437	ACGCAGAACCACCAAGTCAAGCCGGTTTCGTGACTCGGATTCTCAATTCCTCGTTTCATTCCC GAACGAATTTTAAATTCCGAAGAAAACCGCACCGAACACTGAATTTAGATTCTGAACAAGT TTCTTCCGCGAAACAGCACAGCACTTCAATTCACGTGGAACAGAGACAAAGGGAT[G/A]CG TGGTTCGAATTCTCAATCGATTTTCAAATTCGGAACAGCGAACAGTACTTCAATTCACGTC GAAGTAGTCAAAGCGATTCAAATCGATTTCGCGAACTCGTCCGATATTTTCCCTGCACTGAC TTAGTGATTGTTTTCTATCTTTCTCAGCGCTCTTCGATTTTTCCGTTAGTCGAT
>Gm08_4 3903972	GCCCCAGAGAATTCAAGCAAATGACAAATCGTGATTTTTCAGTTACTGTTTTTCTTCCAAA GTGGAAAGCATTTCAGAGAAATAACAGGCAGCATACAAAGTTATTTCAGAAGTAAGCCGCCGC CTCTTCCCATATCAGCATCTCTTGGCCTAGAATGGTGTTCCTCCTCTTGCAGATAA[C/T]TGT GACTTACTGTGAGGCTTTGATGATCGTCCCCTCTCCCACTCATCATCGTGTCTGCTTCCCG GTCTCTGTACCTGTGATGATCTGCATACCTGTCCCTATCTTCTTGTGCCTGTCACGCTCAC GGTAATCCCTTTTCGCGTCCAGATCACGGTCTCGAGATCGTTCACGGTCACGA
>Gm08_4 3960667	GCCACCTTTGGGTGACCTTTTATAATTTTTTTTTGTCTACTTTTTGGTGTTTATGTAACACTA TCACATTGTCATTTTTCAAAATATTATCCATTAAAAAAAGAAATTGTAATCTAATTGTGTAATTG AAACATCTTTGAGGTACTTTTCGTGATTTTGAGGTAGAACTATCGGTTTAA[T/C]GGAGACAT CAAGGCGTTGTTTCAGTTGCCAAATGTGGGACAAGACTTAAGGATACAGTCCTTGACTGGGA GGTACTACATTTTTCATTTGCTTGACGGTAATTTGGTTGAGTCTCAAAAATTATGATTCAATT TGGTTGCAAGTTGTGCCACATGGGTTTTGTTGTTTACTTTGGTATTGA
>Gm08_4 3993352	TTAGAGCCAATGTTGTTGATTTAACTGCAACAGATATTGGTAATCTTCCCTTGCTTTTTCTTA GGTAGAGGCCCTTGGTACACAGGTACATTGATAAGAACCAAAACCTCCACTTAGTAGAAGTTG CAATGCCATTGTTTCATAAATTCTGAGACCCCTTCACTTATGAATACTTGCTTT[G/T]GTAAC ATTCCAAGAAGACAAAACAAATTGAGCAATGCAAGTGTTGCAAGTAAAGTGAGCATTGGGG ATGAGTTTCCAACTCCATAACTTCTTTTTCATATCTTTGGGAAACATTTTCTTCTGCTACCT TTGCAGATATAACAAAGGCTGATTCTGAAAACCCGAAAACTTCAAGATGA
>Gm08_4 3994262	GGACAGGAAATGGAAATAAGCAATTTCAATATAAAATAACCCCAAGTGGTAAAAATGCCCTT ATCTAACAGCTTAAAAAGAGATTTCAGTCCTTAACAACTATACTTGTGCTGCTAAGTGCTAA GAGGGTCTCATATTAATTCATACCTGTGGAAACAAGGGAATGCCTTTGAGGAGAT[T/G]TAG



	TGAAGGGATGATGCAGTAATACAGAGTTGGCCAGGAAAGAAGCACCCATAGATTATAGTAA CAATATCCCATTGGAGGCCTGGACTGATCAATCCATAAGCATACCATGCAGGACTATACTT AGAAAGCACAAATTTGAAAGCCTCCTTCAGACCATCTCTTATGTTGAACTAGTGC
>Gm08_4 5241690	CGCGCCCCCGACGAACACGCCGGGAACGGCGGCCAGCGCCTCCGAGAACCGCCGCGTC GGGCGCGCGATCTCGAACTGCGCGCGGAAGTCGAGGTCCACGAAATACCTGCATGTGGC GGATCCGGTCTGGACCACGTCGATGAACTCGTGGCTTCCGCCGGAGGAGTCCCGCGCGG TCT[C/A]GCACACCGCGGCGTCGTGCCTTCTCTCCCGGAGGAACGCCGCCACGCTCCGCC GGAAGAGCGACGCGTTTCGCTCCCGAAGAAACGCGAGGCTCTCCAACGCCTCCGAAACGT GCGCGATAAGTAAGTTTCTGTAAGGGTCCGCATTGTTGGAGTGTAAGAAAGAGAGACTCGG CGAGTT
>Gm08_4 5620739	TTCTCACCTAGGATACAAACACTAAGGTAGTAAGGTGGTGTATTTATAGATGCCAATAAG AACAAAAATTGCAATAACAATTATAAATTACCAGGTGTTTTACATACAGAAAATGAAGCCAG ACCCAAGAAGAAGTCAGTAGTGAACACTTAAGGCTTAGCCTTTAGTTCTCATAAT[T/A]GTGA TGCTACCTTTTCCCCTTCAATGTGAATCTTGTTTTTCCAACAACGGGCTTCCCAGAAAAAT GCTGATTTTGGGTAAGCTGATGGATCTTCCATACTTTGTTGTTGGGACTTCAAAGGAGGTG GTGGAGACAGAATGCGCTGATTTATTTCTTCAGAGTGACATCTCCATGGGCA
>Gm08_4 5632650	GAAGAACAACAAAAAGCTTCATTTGCCAGCATGAGTTTATTCTTCTGTTCTTGAAAAACAAGG GTCTAAGTGGACAATTAACAGTAAGCAGAGAGGATTACAAATTTGTCTGGACATAGTGGTT TCTGATTTACAATTGCCTATTTGTTGCTTCTGCCACCTCAAATACTACTGTACT[C/G]ATGAC TAGGAAGATTATAGACAAAAAGCATCAATTTGTTTGTGATTGTAATTTGCAATTTAACTTGA AAAAGTATTTAGATCTAGATTGTGGCACAATGCATTTTGATCTTTACTCTTCCCTTTCTTCA TTAACTTGGGTAGTATTAATACAGATAAAGATCGTTCTTGATGAACAC
>Gm08_4 5819046	CAAATATAGATTTGTTTCATGACACAAATCTTTCCTACTATACAGTTAAAAGCACACCACTCGGA ATCATGTCTATTCGTAACACTTTTATCAAATGTATGCTGATATAGATGCATACCGTATATCCTA TTATAAAAAGCAACACCACTCGACTATATAGTAGTTAGGTGACACATTTTCCCT[G/A]JAGAAA CTTGGAAGTGAAGAACGTGATGCTCAGAACTCCCTTTCTGGAAACATGATAGGAGCTACGA GAGACCACAAGTACAGCAGAGCAGTTGCCAGGAAGTGATGATTGCAACCCACACAGAAG GCCACCCACGTCAACCAACTTCCCGCTCTCTCCAACAGATGTAGACCAACCAG
>Gm08_4 5828129	GAACCCGCACGTTCCCCGTTAAAGCTTGATATCACAACATAGGGGCAACACAAGAAACCAT ATATAAAGAACTTAATAAGATGACATAAAATTCTAGAGCAAAAAAATTAACAACAGTTCAA GATTATCATATCTAGCGTCCAAAATGACCAGCAAACTGAAATGTAGTTCAATTC[G/T]GTAT ACACCTCAAACCAAACTTGAAACACTAACTAGATACTACTTAATACTTAACAGGAGCAAGG ATTTCAATTTGGGAACCAAGCTTCTCCTCTGCAACCTTCTCAGCTTTCACCCTCAATTTGGC AAGTTGCTTCTTCTCTCATAACACAATTGTGCTCTCTCCTTCTTTCTTC
>Gm08_4 5831798	CAACACCCTAATAAATTTTATTACCACTTCAGCATCTAAAAGATTGAACAACCAAAATTC AACAAAAGTAGAAAGAAAACTCACTTGAGAGCATCAGGAACAACCATCCTCTTGGTCTTGT CATAAGGAGGAGGAATGCCCTCATACACCTTCAATCGCGCAAGAGCAGCTTCCC[C/T]GCG CTTTGTCTTGTGTGATCATCCTGCAAAACACAAACATCCGTCTTAGAAAACCTTCCACAAC AACACCAGTAATCTTCAATTACAAGTTTATAATGTCAACCATAAACACATTAGCCCCAAATAAT GCAAAAAACACTACACATTAGCCCCAACATCAATAAAATCCGTCCCACTAGA
>Gm08_4 5844445	GACCGTTCATTTGAACTGTTTGACGAGATGGAGCGGATCAGGTTGATCCGACCCAATCAT TATGTGCGGCCAAACAAAAAATATATTATATCTTTAATTTGCACTTTTACCAAATTGACGACTT GTGTAATACCTAGTCTCAGTGCCCAAAACAAAACCCCTTGCCTCCTCGCCGCACA[G/A]TTA CACTCCGCAGCACACTGCAATCGCCACCACGGACCAAGCTAGTTCCGCCGTCGACGTCGT CGGCCACCGTCACGTCTCGCCGTCAGGTGCCTCCCTCTGTTTCTTTGGGGTTCTGGGCAT CTTTGTTTATTTGCCACGAATCTCTCTTTACCAACTCTTAACTAAAACCTTGCTGATT
>Gm08_4 5905582	CTTTGTACCTCAGGCACACAGGAGTTGTTGTTTATGGTAATGAGTATTACTTTGGTGGGGT ATTCAACACTCTCCTGCTGGATCAACTCCATATGGAACCTCCACTAAGAGTGGTAGACTTAG GTGTGACACATGTTCCCAAGGATGTCTTTGAAATGTATTTGCAGGAAATCAGTCCG[G/A]AG TACCTGCCTGAAACATATAGTCTGCTTACTCATAATTGCAACAACCTTCAGCAATGAGGTTGC TCAATTTTGGTTGGTGGTCCATTCTGAATATATTCTCCAGCTGCCCAATGAAGTCATGA GCAGCCCAATGGGAGCTCTTATATGTGAGTTATTGCATTTCCATGTCACTTACC
>Gm08_4 5965493	TTAACTGCACTACACAAAACGTACCAGTTATCTCACTTCCATCTGAGAGACTGTCTAATGC CTGTATAAATGAGTCCTAAAGACCTCCCAATCAAGCTAACTAAAAAGAATGAGATTGCATCA CTCACTTTATCAATAACTGCTCTAATTAGAGGCATGGTGATGTCAGACAACCTCAA[G/C]AAAG AGGCTGAAATAATATCGCCTGCAATGCATAAATTAACAGCAATTATTACATTAATTGGATGA AAATTTCTCGAAAATGAGTTTGATTCTATAGTGAGCATGGGTGCGAGAGGGTACCTTGATT

	GATTTAAAGTATCATGATCTTAACCAGCTAAAACCTTAAGAGTAAATATACCC
>Gm09_5 927324	TCTTGATATGTTTTATCGACTACTTTGGATTTATTTCTGCCATCTGAATATGTCTGTTTCAGTAT ATTTAAGGTGCTCAATTCTTTCAACTGCACCATGTCCTTAGGTGATACATAATTAAGCAAGTT AATTTACTCTCAACTTTTTCCAATACATGGTGTGAATGTTGCTAGTCTCTT[T/C]CCAGGGA GGCACTTTTCACCCCTTTCTTGCAAGTCTGCTGGGTCTATGTTCTTTGACTTATCTTGCTCCA AATGCTGAGTCTTGGAAGAATCTCTCTTTCAATTGAACCTTTCTTGTTGTTTTGTTGGAGGTG GTTTAATCTGAGGTCTCTTATCCTGTTTTAAGACAAGTTCATGTGAAC
>Gm09_1 9158732	GTTATTGATGTTGGTGTGTTGGTGTGCTGTTGTTATTTTCGGCCGACTCAGAATCCTCG GCCGAGACAACAACAAATTGTTATTGTTATTGCTGCTGTTGTTGTTCCGCCGGCAAATTC GATGCCAAAGAGGCGAATGCCTGACTTCTCTCTGACAAGAGGAGCCGGGCGGATGA[T/C]G GGGAGTTGGGAGAACGAGTCGACGTTTCATGAAGTCGTGAGTTTCTCTCTCGGAGGTTGTT GTTGTTGATGATGTCTTGTCATGAGAGGGACTAATGGGAGCTAGGGAGAGGTTTGTGAAT TGTGATGTGAGGGTGTGTATATAGAGAAAGCTAGGGGAGTATAGTGTATGAAGGTGTG
>Gm09_3 0684887	GACGGAGTTTGGCCCTCGAGTGCAAAGGCAGAAGGGAGCTTGATTGCAAGACCCACCCGT CGATTATGGACAAAAGTCGGTCTTAATGATCTGACGATGCCGAGTGGAAGGGCCATCGCTC AATGGATAAAAGTTACTCTAGGGATAACAGACTGATCTTCCCTAAGATCTCACATGGA[G/C] GGGAAGGTTTGGCACCTCGATGTCGGCTCTTCGCCACCTGAGGCTGTAGTATGTTTCCAAG GGTTTGGTTGTTTCGCCATTAAAGCGGTACGTGAGCTGGGTTTGAACGTCATAAGATAGT TCGGTCCATATCCAATGTGGGCTTTAGAGCATTGAGAGGACCTTCCCTAGTATGAGAG
>Gm09_3 0684888	ACGGAGTTTGGCCCTCGAGTGCAAAGGCAGAAGGGAGCTTGATTGCAAGACCCACCCGTC GATTATGGACAAAAGTCGGTCTTAATGATCTGACGATGCCGAGTGGAAGGGCCATCGCTCA ATGGATAAAAGTTACTCTAGGGATAACAGACTGATCTTCCCTAAGATCTCACATGGAT[T/G]G GAAGGTTTGGCACCTCGATGTCGGCTCTTCGCCACCTGAGGCTGTAGTATGTTCCAAGGGT TTGTTGTTTCGCCATTAAAGCGGTACGTGAGCTGGGTTTGAACGTCATAAGATAGTTTCG GTCCATATCCAATGTGGGCTTTAGAGCATTGAGAGGACCTTCCCTAGTATGAGAGG
>Gm09_3 9232035	GAGGCGCCGCCGAGGACCCCTCCCTCGCCGGCCTCCACGCCGGCGGAGGATGACCTCG GCGAGATTGTGAGCTGCCGCCGCTGGGAACGAGTTTCGACTCGCCCATCGACCCGATGA GTACCGAGTTCGTGTTCTCGGACATGGACGACGGGTGGCCCTATTCTCACCCATGGTACCA [A/C]GGCATTATGATGGTGGGTAAGTATTGATTAGCGACATCAACAACATGGTTTCAATGCAAG AGTCAGAGAGCATGCTTCTTTCTTGTTGGCCTTCACTTAAGCCAATGCTATACTCATTCCA GTATGATTATTTTTGCGTCAGTGATATGATTAACAGATCATCATGCTTTGATCTCTTTCT
>Gm09_4 2479372	AACTGATCTGTATTTCATTGTTTTCTACTAGCATTTTCAGAATTAATGACAGCATTAGCCACT GAGACTTTGAACCTTTGTAGACTCATAACTTTCTCTTCCAAAGTACCACGCATTATTAGCCG ATGGACATTAACCTACTTTTTCTGACCTAACCTGTGTGCTCTATCCATAGCCTG[G/T]AGGTT ATTGGCATTACAAAAACAAATATTAGATATTCATTCCCTTAAATGCAAGATGAGAAAAGCTT CATGATGTTAGGCCTCAAAATATAGAAAGGGCATGCAAGAAAACAGTATACGCACAACACAA GATGTTAGCTTGAGCTCTAGAAAAAAGTGATTACAAGGTTGTGAAAAAAGA
>Gm09_4 6356127	ACCGGGAGGTTTCGCGGTGTTCCATTTGCGGATCCCTAAAGATACGTATAGTTTCAATTTCGT TCCTTGATTGTTCAATTGTTTCAAAGTAAATGATTGCTTGAAGGCTTTGAAAGCTTTGATTCTT CGATGCCGTATATACGTGAAATTTGTTTTGAAGAAAAGAGTGTGGTGATGGTGCTG[A/G]GAAA TTCGATTTTCGATGATTCAAATGGAAGTGTATGTTTCTGTTATAAGCATGATTTGGTTAAG ATAGCTGAGCCATTGATTGAGGAGGAGCCATTGAAGCCTTGATTACATTGAAAAGGAATG TGAGGAACATAAAGTACCTTCCTTCAATGGCTGATATAAGCTTTAAGAATAG
>Gm10_4 227818	AAGAAGGTTGGATCTTTCAGCATGCCCTCCAAGTCATGGCAATTGGAGCATTCTGATTTCGT CGGAGAAATGTGAAAATTCCTGTGGATCATCCTCAATTAGTTCACTTTCTAAACATTCTGAA GCAAACGAAATAAAGCATGATGCAAAAATTCAGCTTTGCCAGAAAATGTAGATCA[T/C]AG TGACAATGTTTCAGAAGCCTCTACACTTATTCAGACCCACAATAGTCCATCCAGTACAAAAG ATTGTGATTCTCAGAATACTGATAATTTCACTACTAATTCAGAACAGGGTGTCTTCCCCAC AAATTCTGAGACAGAATCCGAATCTGAGCTCTGGGAATTTGAAGAATGATGAA
>Gm10_6 376650	AGGACTGCATTTCAACTTGGCAAAAATATGAAGGGGAAAAAATACGCTGAATAAATGATG GCAAAACCCCCCTTATAGGGGAAAAATCATGAATATTCATGACACCTAAACAACATGCAGTT TAATATCATATGATCTACAATTGATCATTCTATTATAATCAGATGCCAGGGAAAGA[T/C]AACA AGTGTAATTTGTAAGTATTAAGAATCTTCTTCTTCCGTTTTGAAGCACTTCCATCACTCTCA TTTTTCCGCTTCGATTCTCGGCGCTCCCTCTTCACTCCTTTTGCTTTGTAAGGTAAAGGAGT TTCTCCAACCTAAACAATCACAAATTGTTGTAAGTGGTTAGATATCCTTCA
>Gm10_3 9929704	GAAGTTTCCGAGCCCTTCGGGTCTATAAGAGTCAAATCTATCTGTTGCAACAACGACAACG ACAAAAACACGATGGCGATTTCAGAAGCTGAATTGAAGCTGTTAAGTGTTACAAAACATGGAT

	ACATACACCAAGATCGCTTTTCTGGGACATCTGGAGCACTTAACACAATTCACCC[G/C]CA GAATGTCACCTTCTGTATCAACTTGCACATAATAATCCTTCGGAGGAGGCCAAGCCCAAG TTTCGTAAAATACAACTAAACCAATAAACAACAACATAACATAAGCATGAAATTAAGATAA TCACTGCAAGCTTCTTCGTCATTCATATTTTAAAAATCATAATAATAATTAAA
>Gm10_4 2374383	ACATTATCATTTTCAACATTATTACTGCTCTATTGAGTTTGTAAAAGAATATTAATAGATGGC AACCTCAAAGGTCTTAGGGGCTGCTTTCATTGTTTTGCTCATTGTGGACCTTGCGTTTGCTG CTAGGGTTACAGATAGGTTACTTGGTGGGAGGGGTGGCGGTGGTGGTGGCGGCG[T/G]TG GTGGCGGAGGAGGTGGCGGTGGTGGTAGTCTTGGACGTGGTTCCGGGTATGGATCAGGG TATGGCTCAGGTGGGGGTGAAGGGTATGGTGGAGGATCACTAGGAGGAAGCGGCGGTGG TGGTGGTGGCGGAAGAGGAGGAGGTGGCGGTGGCGGTGGTGGGACCGGCGGTCTGGG TA
>Gm10_4 2374384	CATTATCATTTTCAACATTATTACTGCTCTATTGAGTTTGTAAAAGAATATTAATAGATGGCA ACCTCAAAGGTCTTAGGGGCTGCTTTCATTGTTTTGCTCATTGTGGACCTTGCGTTTGCTGC TAGGGTTACAGATAGGTTACTTGGTGGGAGGGGTGGCGGTGGTGGTGGCGGCG[A/C]GG TGGCGGAGGAGGTGGCGGTGGTGGTAGTCTTGGACGTGGTTCCGGGTATGGATCAGGGT ATGGCTCAGGTGGGGGTGAAGGGTATGGTGGAGGATCACTAGGAGGAAGCGGCGGTGGT GGTGGTGGCGGAAGAGGAGGAGGTGGCGGTGGCGGTGGTGGGACCGGCGGTCTGGGT AC
>Gm10_5 0508726	ATGGGAGTGCATTAGGTGCTAAAGAAGTGGATGCTACAAGGAAGAATCTTGGATGGCCATA CGAGCCTTTCCATGTGCCAGAAGATGTCAAGAAGTATGATGGGGAGTTAGAGATATCTTTG TTTCCTCTTAGTCTGCTGAGATTTTCATGATCTAGAATGAACTTTCTTTATTTTCC[G/T]TAG GCATTGGAGTCGCCATACACCTGAGGGTGTAAACTTGAAGCTGAGTGGAATGCCAAGTTT GTGGAATATGAGAAGCAATACAGTGAGGAAGCTGCAGAGCTGAAGGCTATTATTACTGGCG AATTACCAGCTGGTTGGGAGAAAGCACTTCCGGTGAGTAACTAACAACTTCTA
>Gm11_8 599758	GTCAGGTTTTGTGATTTTGTAAACAAGATCTGATTCTCATTTTGTACTACCGACATATGAGTA GTTGGAAGAAGGTCAAGTGAAAATGCAGTCAAGATTGGAAGAAGAGGAAGAAGCGAAGGT TGCTCTTATGAGTCGGATCCAGAAGCTAACCAAGCTCATTCTGGTTTCATCAAAGA[T/C]CG CAATTCCAGGATATCTAACTGATGCTTCCAGCCATCAGCAGAGTCCTTCTGTTGGTGAGGA TGATGTGAGTTCTTCTTCAATTGGTGTGAAGTGTGTTTTACCGTCTTTTTAAATAATATTC GTTCTAGATGTGCAAGTACTTTGAACCTCCTTATCATGATTAGCTGCAATATTAT
>Gm11_1 5759081	GGGCACCCCACTGATATCATCCAACATGGGTGAATGCTCAGCAAACACCCCTTTCTTCACC TTCTTGACAAAGGCAATGCAAGCAAGGTACAAATACTCATTAGAGAAATTCTCCAAAATATC CTCATTGTGAATGGACTTGGGCTTCATATACTTGTGATCAATCAACTGCGAAGACC[T/C]AAA TATGAAGGGCAAAAAATGATAATCATCAAGCCCCCACACACCATGGGACCCAACAGGCTCC AAGCAGTACTGTAATTGCAGCTTCCCTCATGAGGTCAANNNNNNNNNNNNNNNNNNNNNNNN NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
>Gm12_5 642827	GGAGGAAACCAAGTCCAAACCTACGTTCAATTGCGCGGCGAGTTCGGTCAGAGCAATCTG ACTACATTCTGAAGCCTCTCGTTAATCTCTTCTGCAATCGAGTCCCAATAAGACTCCGACA TAATTTCTCCCTGAGTCAACATCAGTTCTCCGTGTTCCGTTACAACGCTCTGAGCCT[T/C]TT TCTCCACGTAGTACAAGTCAACTCCAGTAGCGTCAGCGAGATCGATCAAAGAAATGCGCCC TAATTTCTTCACCTCCGCCACCATCTCGTTCTCAGCTGATCCTAACAGCATGCACGATCGC ATTAATAAACGTTGACTAAGAATCAGCACAACATAGTAAACGAAGCACGGAAAC
>Gm12_9 267296	GTCGTGCCGTGGTCCACGCGCCGCCGCCGCCGCCGCTCGAGCTCCTCCGGCAGCGA ATGCTCGGAGTGCAAAAACAACGTTGCTTCTTTTCCGCCTCGGAACAGGAAACGACAACA TCCTCGTTGAAACGACAGCGCGAGGACGACGACGATGCTTCTCCGATCCTGATCCCAG[T/ C]TTTGTTTTATTTTATTTTGTTCGAACGAGGAATAATTATTTATTTACTTATTTTAGTT TGTCGCTTGAAATCAGGGTGTGAGAAGGGTTGGCGTCTCAACGGAGGTTGCGTGTGAC GGAGAATTGTCGTTGAAGGAAAGAGTGGTGTGGCATACGACGTCGTTTCGTGATGG
>Gm12_1 4154520	AACTTAACTTCCATTCCAGTATAAGCAGATAGGCAGATACATCGTATTCACGTGGGTCTGCA TGCCCCAACAACAACTCTCCTTCCACTATCCATTAATACCATCCCCCTCCTTACCCTAATTTA ACATCACACATCCAATACATTTGTTTTACCCACGCTCCCCCTCTCTCTCTC[G/C]JACACT ATCATACTACCACTCAGCTCAAAAAAAAAAAAAAAAAAATATATATATATATATATATATC TTTTCTTTTAAATTTTATAATTTTGGTGAAGATAGCAAGAGAGAAAGGGTTGGAATTGGAAT ATCTTACCGGAGAATGGGCGACGGCAGCATCGTGGGCCACCTGCCGGA
>Gm12_1 4154521	ACTTAACTTCCATTCCAGTATAAGCAGATAGGCAGATACATCGTATTCACGTGGGTCTGCAT GCCCTAACAAACCTCTCCTTCCACTATCCATTAATACCATCCCCCTCCTTACCCTAATTTAA CATCACACATCCAATACATTTGTTTTACCCACGCTCCCCCTCTCTCTCTCT[T/G]CACTA

	TCATACTACCACTCAGCTCAAAAAAAAAAAAAAAAAAATATATATATATATATATATCT TTTCTTTTAAATTTTATAATTTTGGTGAAAATAGCAAGAGAGAAAGGTTGGAATTGGAATA TCTTACCGGAGAATGGGCGACGGCAGCATCGTGGGCCACCTGCCGGAG
>Gm12_1 4158308	ATCCAGGATGTTAAACCGAACACACTGTATAACTTACCGGCATATGTCTCGCAACATCTTTA GTCTTCTCCGCCAGCTAAGCTTCTTCTTCTGACCGCTCACATGAATCAAGTAAACAAAGAT CCCATCTCCATGTATTCAAGTCACCATTGACAAGCGTGGGGGCTTTGTGCATGCAC[A/C]AAG AAATAAGATAACTGCATAAAATAAGATAACAGATGCTATCAAAGAAATCATGCTACATGAGA TATTCTTACGACAATATATGCACAATATCATTCTTAGAATTTCAAGCAAGGAAGAATTTGGTA ATAATTCCAGCTGTCAAAAATATCTAGAAACAAAATGTGCAAGATATCAATA
>Gm12_1 5498752	TAACCTATTGTTTTTCCCTAGATTTCAATTTTGTAAAGAAATATAAACTGTTTTGCACTTTA CAGGTTCCAGGAAATTTGATCATCTCAGCTAGATCAGACGCCATTCTTTGATGCTTCTCA AATGAACATGTCACATGTCATAAACAACCTTGCTTTTGGAAAGAAAGTACTC[G/T]TAGGGC CATGAGTGATGTGAAGCTCTTGATACCATACATAGGTAGCAGCCATGACAGGCTCAACGGC CGGTCATTCATCAATACTCGTGATTTAGGTGCCAATGTTACTGTAAGTTCAATATTTTATATG AAAACTTATAATTTAGTGTTAATTTTACTATTTGGCTTTTGTGAGTTGG
>Gm12_2 3879822	TCGATCGACCTCTCCATCAACACTGACCCCACTCCGACGAAGACGCCGTCAGCGCCGTC AACAGTAACGACCACGCCAACGCCGCAACAGCAAACGCAAGCTCGAGGACGACTTCGAC GCCTTCACGGCGTCGAAGGCCGCGGATCTCGCGAAGCCGTTGAGAGAATCGCAAATTC[A /G]GTTCAAGATCATATTGTGAAGGATCACGACATGAAGGAGAGGCTGTGCTTGGAGGTGGA GAGGCTTGGGTTGAGCGCGGTGATTATGGGGAGTAGAGGGTTTGGCGCTGTGAGGCGTG GCAGCGATGGGAGGCTTGGGAGTGTTAGTGATTACTGCGTTCATCACTGTGTTTGTCTGT T
>Gm12_2 4557725	AGTGGCATAGACCCAATTTGGGGCAGTCTCAGGTGTAACAGCCACCGGATACTTAATGCTT GAAGGATTCACCGACACATTTAGGACCGAGCTGTTGACATGAAGGTATTTCTATCATTAC CCAAGTCCTACCCAGGGTGTCATTTTGGGCAGTAATCAAAGGACCCCATGTTCA[G/A]GC GATAAACAGTTTCAAAGGCAAGTTCAGAAAGTCCACTGAATGTTGCTGGCGGGTTAACAGC CAGTGATGATCAAAAAACAACCTCATTGCGCATTGACACGACTTCAATTGCATTGACAAAGG CCACTGAACCATTGGAAGGAATGAAGGTGACAACCAATGTATCAGAGGTAACATT
>Gm12_3 2742628	ACAATAAGCAAAAATCTTCAGAAACATTTAATCAAGCATCTTGAGGCAGAAATTGAATAGT GCTAAACATGGAGAACTTACAAAACCTCATTGAGGAGATCACTTGACATTGAACTCCCT ACCTACACCAACCTTAACCGTCTCGTTTCCCAGGTGATTCATCCCTCACTGCCT[G/A]GTT GAGGTTTGATGGAGCACTAAACGTTGATGTGACAGTTCCAAACCAACTTGGTTCCTTATCCT AGGATCCATTTATGCTTTCCTCATACGCCCTGTTATCACTGCTGAGAAAGCATACCATGA GCAGCTTTCGTTGCTGAGATCACCAACAGTGCTTTGAGCCATCCTCCATGA
>Gm12_3 8201850	TTGCAATGCTGCTGTTTACTTGCCAACCTGCTAAGGAACCTACCTTCACTGCTGAGGGCTTTG AAGTGTGTTGGGACAAATCATCTGGGGCATTTCTCCTCTCGCGCTGTTGCTTGAGGA CTTGGAATAATCCGATTACCTTCAAAGCGCTTGATCATCGTTGGTTCAATAACAG[T/C]TCT AGTTCATTAGCATGTTTATTGTGAGTTCAGTTCCTATCACACCATTAGGTTGCATTTCTTAT GCTGATTGTGTGATTCAACAGGGAACACAAACACATTGGCTGGTAATGTACCTCCCAAGG CTAACCTTGGTGACTTGAGGGGACTTCAGGGTGGTTGAATGGGCTAAACAGC
>Gm13_1 275756	TTTCCTTAAATTTAATGAGTCAGATTTGAGAAAGTTGACATAAGCTTAAGAAGCACGGCCA AGATATTAGGGACAGTTTGTGCGTGGCTGGAGCCTTAACCATGGCATTGGTCAAAGGACA GAAGCTGTTACACACGGAGTTCCTTCTTCTATACATCTACCGGCAGTCAAGGTG[T/C]TG ATTGGCTGCTGGGTTGCCTATTGCTCTTAGCAAGCAGTGTTTTTGGTCATGTTGGATGATC CTGCAGGTAAATAAAAGCCGATTTATTTTCTCATCCTTCTTTGATACAATCGATGAATAGAA AATTTATTATGTCATATGAAATTATCACATAATTATAATTCAAACAAATATCT
>Gm13_9 272222	ATGAATGAATATGCATAAACAATTATTCTCTTGATATTTGAGTAACGATTATTTTATTATT TTGTCAGAAACAAGGCACAACTGGTTGTGATTGCTCATGACGTGGACCCAATTGAGCTGGTG GTCTGGCTTCCAACATTGTGTAGGAAGATGGAATTCCATACTGCATTGTCAAG[T/C]GCTA CTCTAACAATCATGAGCATTGTGCAGCAAGATGTTTGAAAGGGCAGTCAAACAGGAGAATT ATTTCTACGGATGAAGTTAGAAAACACCAGATCTGGCCAAATGTGGACTGTATTAAGGCA GTGTTTATAAAATATCTCAGCATATGAAGTTTACCTTGAAGGTACTGCTATA
>Gm13_9 272223	TGAATGAATATGCATAAACAATTATTCTCTTGATATTTGAGTAACGATTATTTTATTATT TGTCAGAAACAAGGCACAACTGGTTGTGATTGCTCATGACGTGGACCCAATTGAGCTGGTGG TCTGGCTTCCAACATTGTGTAGGAAGATGGAATTCCATACTGCATTGTCAAGA[G/C]CTAC TCTAACAATCATGAGCATTGTGCAGCAAGATGTTTGAAAGGGCAGTCAAACAGGAGAATTA TTTCTACGGATGAAGTTAGAAAACACCAGATCTGGCCAAATGTGGACTGTATTAAGGCA

	GTGTTTATAAAATATCTCAGCATATGAAGTTTCACCTTGAAGGTAAGTACTGCTATAA
>Gm13_2 1371345	AGACTGATTGCAAACCTCAATCAGAGCTTTCACCTATTGTCCAAATACTGCACAAACAAATT GAGTCAAGTGCAAACAAATATTGGGAGGTGAGATGAAACCAAATTACATTTCTATTTCTTTT CTACAAAACCTGTTAGTACAATCACAAAATTTCCATAAAAAGATTAATACAAAAAA[T/G]TGCCA CCATATTTCTTCTTTCCAAAGCCAGCTAAAACGACAACGGAATACCCTGGAGTTTATATCT GAGAATCCTAACAACGGCATTATAAGACAGTTATGGTAAAAGAGCTTTTCAGTTGTGCTTGGC ATAAATCACCAAGAGCTCCCCGAACGTCCATCTGATCACACAGGATTATCA
>Gm13_3 1847248	CCTCTGGCAAATTCTCTACCAAAACAGTATATAACTAGAGTCCAAGAATAAGATGCAGGATA ACAGAGAACAACCAAAAAAGAGTATAAAGCTCCTTAAGTAAATACCTTATGATCTCTAGAT TCATTGAAACGAAGTGGATTGAGCCGCTTGACCCTTCTCCCATCACCACTAACAA[A/C]CTG CAGTCCCCATGGGAGAATTTTTTGCTTAGGACTTGAAGTATAGATTGCAATGCAGTTCATGT TGACAAACAAGGAAGGAAGAATCATACAAGCAGTGAAGACTCCTTCAGTGCAGCAACAATA AATGCATGGTCTCGCGTGAGCTTCTTTATTTTTCTAAAAGAAGCAATTACTGAT
>Gm13_3 5524268	AGTGATCAAATGAATAGTGAAACTCTGATTCTGATTGGAGAGTGATACCAAAAGTAATTA GAAACTTCTGGCTCTAAGTATTGTGCTTTTTTATTTGTGAGGCATTTTGGTTATTGCATTAT GTGTTATGCTGTAATGCATGCAGGCTGCAAGGAATGCTGGCGTGCCAGTAGTGT[G/T]TGAT GCGGGGGGCATGGATGGGCCCCCTCCACCACAATTATTGAATTTTGTGATATTCTGAGTC CTAATGAAACTGAACTTGCTCGCCTTACCGGAATGCCAACAGAAAGTTTTGAAGAGATTGC ACAGGCTGCTTTGAAATGCCATGAATTGGTCAGTTCAAGTAATGACTTTAGGGG
>Gm13_3 5524361	TTATTTGTGAGGCATTTTGGTTATTGCATTATGTGTTATGCTGTAATGCATGCAGGCTGCAA GGAATGCTGGCGTGCCAGTAGTGTGTTGATGCGGGGGGCATGGATGGGCCCTTCCACCAC AATTATTGAATTTTGTGATATTCTGAGTCCTAATGAACTGAACTTGCTCGCCTTA[A/G]CG GAATGCCAACAGAAAGTTTTGAAGAGATTGCACAGGCTGCTTTGAAATGCCATGAATTGGT CAGTTCAAGTAATGACTTTAGGGGTCACAATGAAATGATTCTGCACACTTTTCTACTTGAAT GCCATATTCATTGCTTATGGATAGACAAGGTTTTAGATTGCGGTTATAGTTGCAA
>Gm13_3 5526193	ACATAATTAACCTTCATTAGATGCATAATGCATTCCCTAGTACAATTCATGAGTCCATTTATTA AAATTTCTCTAGTCTGTTGATCCTGGTCTGCGAACATAGGAGAATCTGTTAGTTCACCTCCC TATTATTGGTTGTTTTACACAGGAATAACAATCAGTGAAGATTTTGAACGCA[G/A]TAGGA ACAGGAACAGAATAACTTATGGCAGAGAGAAAGACCCTGGAAAATGGCAGCAAGTAAAGCA AAAGCCCTTTTACAGAGCATTCTGCAAGAAGGATTCTTCAGCATAATAGCTTGTTGTCATT GTAATAAGATTAGGCCAAGCACTTACCCTTTGGCAATAGAAAGAAAATGAAG
>Gm13_3 5526194	CATAATTAACCTTCATTAGATGCATAATGCATTCCCTAGTACAATTCATGAGTCCATTTATTA AATTTCTCTAGTCTGTTGATCCTGGTCTGCGAACATAGGAGAATCTGTTAGTTCACCTCCCT ATTATTGGTTGTTTTACACAGGAATAACAATCAGTGAAGATTTTGAACGCA[G/A]AGGAA CAGGAACAGAATAACTTATGGCAGAGAGAAAGACCCTGGAAAATGGCAGCAAGTAAAGCAA AAGCCCTTTTACAGAGCATTCTGCAAGAAGGATTCTTCAGCATAATAGCTTGTTGTCATTG TAATAAGATTAGGCCAAGCACTTACCCTTTGGCAATAGAAAGAAAATGAAGA
>Gm13_3 5582717	ATATGATAATATAACTCTACTGATACTTCAATATTAGAAGTGAAGTACATGATGGATT AAGTATTACGATACACACTTATCCCTAACAAATATTGCCTATGGAACGAACTAACTGTA GTTTCTAAAGAATTTGAGCATTGGGAAAGCACAAGGGGAGCTTCAACCTGCATC[T/C]TCAT TTGATTAAACTTGAGCGCTAATATGGGTTTTGATAACATCACACAAGTGCTGAAATCTGTG CATCATTTTTGTGATCTTTAGCTACATGGCTACTGCTTCATAATCTTCTAATGACAGTTTCA TGTCATCATTTGTTGAAGGGAATGCTCCTGAAACCTTCAAGGGCTTCAATG
>Gm13_3 5582814	TATTGCCTATGGAACAGAACTAACTGTAGTTTCTAAAGAATTTGAGCATTGGGAAAGCACAA GGGGAGCTTCAACCTGCATCATCATTTGATTAACTTGAGCGCTAATATGGGTTTTGATAAC ATCACACAAGTGCTGAAATTCTGTGCATCATTTTTGTGATCTTTAGCTACATGGC[C/T]ACTG CTTCATAATCTTCTAATGACAGTTCAAGTGTGCATCATTTGTTGAAGGGAATGCTCCTGAAAC CTTCAAGGGCTTCAATGCTTGCAACCCTATTCAATATTTAGATTGATAAGATACTTTAGACTAC TTTCAGGCTCATATGTCAAAAAACATAATATTGATTAATTTCTTTGTGCCT
>Gm13_3 5600102	TAATAATGTAAAACTGGCCATCTGACTACAACACTACAAGTACCAGTACCAGTACTTGTA CCAGATATATGGTAACTATGATATAGACAATTTCAAGTTATTCATTATCACAACTTCTGGTCT CTCTCAAAGATGCAACGGTAATTGCTCTCCTATAAATTGAATCAAACAGACAGG[G/A]ATGTT GCCAACTCCAGTTTCAGACCACATCTAGCAGCAAAAGGTCATTTCTAACTCTGATTCTGTTT TGTGCAGGAACAAGAAGTTGTCACTGTAAGAATAGTCTCAAAGATTAATGAAGTTTGCCTTT TCTTATTAGCATTAAACAGATAGACATTGAGATTTTAGATGTAATGTTGATT
>Gm13_3 5617464	GTGATAATTGCTGCACGGAGCAACAACGAGATGGATGCATCCAACTGAAGAACGAGTTCC CTGAGCTTCTTTCCATCAAGGAATCACTGATCAAGTATGTCTTTGAGCCAAACAAGAAAAC

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>Gm13_3 5618692	ACATCATAATAATAGCAGTACAAAAATAAAAACTTACATCATAATTAACCTCGATTACTGATT GATGTTAAAAAGAATAAGTTATATCATAACTAATTTAAAAAAAATATTTTTACGATAATTAAT TTGATTATTATATTGTCTAAAAAGGTTAGGATGTGAAGTTATGATTATGGGG[A/T]GCCCCCG TAGAATAAAAGCAAATCGTGTACAAGAGAGTAAATTTGGCAAGTTGCAATGTTGGCAATATA GGGAATTTCTGAAGCTGGGTAAGTCTCAGACATTGGCTAGTTCCTTGAGAAATGCAACAAC TTCATCTGGTGATCCAAGCGTATACCGAGCATTGGTCTGTGTTCTGTCCTCAA
>Gm13_3 5690746	ATTTCTATTGTGCTAGGTGAATTTAACCCTAATTTTATTCTGAATTTTCAGCCTCCTTAATTCT TTATCTATTCATTTGTAACCTAAACCAAGAACATCCTAATTTTGGTCCCTTTTGTGAAATATA GATGGCATGCTATTGCTTCTTGGACTTGGGATGCACAGGACGAAACGTGTG[A/G]GATTTGT AGGATGGCCTTTGATGGATGTTGTCTGACTGTAACTTCCTGGAGACGACTGCCCGCTGA GTAAGCTACTGATAATTTTACATGGCGATAGACATTGTTATTTTACTTGAATCCATTTGTCTT GGTTAAATTTCTTGATTGATCATTGTGTTTTGGGAATTTTCCAGTAAAG
>Gm13_3 5759476	ATTGTCCATAGAAAGCAAAGTTTTATTTTGCACGATATAATAGGCGCCAGCGTCATTATCCA TAAAGTTACCAGTATTGATATATGCCCAATTTTTTGTCCACTTCGGTGAATCTTGCTGGAC CTCCAGTATCTGAATCATCATCATATGTGCTTCCATTGTCTGTTACTATCTTTT[C/A]ACCACA ATTTATATGAAGAGAATATAACGCTGTGATAAAAAATTAATAAAGTTAGCAATAATTTAAA ATATTATTTAGAAAAATACATTCCCTTCAAATTTGATTTTTCTATTCTTTTCTTTTCTACTCAA ATAAAGATGCAGTGGAGTTATGTTAACTTATATAGGTCTAACTAGT
>Gm13_3 5789590	TTTGCCCACTCTCTTTGTTCTCTCTTCTTGTAGCCTCTGTTCCGTTTATTCTGCTTGCCTT TTTTTTTTTTTTCTTTTCTTTTTCATTCAATTTCTGAACTCTCCTTTTCAAACCAAGATACG GTGGTTCTTTTTCGGATCCTGACAAACCGCCGACCATGATGCGGTTTGTGAT[T/C]CCTTCA AGCAGCATACGTGTGTCGACTATGCTCTTCATTTTTATGTTCTGGCTTCCCTACACTTTGG ATCCAACGCTCAACTCATACCACAAGATGAAGGTACTTAATTCATTTCCATTTTGTTTTTT TTTGTATGATTTTGTATGTTACTATCTCTGTTTCAAAACACTTTTGT
>Gm13_3 5789620	TTTAGCCTCTGTTCCGTTTATTCTGCTTGCCTTTTTTTTTTTTTCTTTTCTTTTTCATTCAAT TCTGAAACTCTCCTTTTCAAACCAAGATACGGTGGTCTTTTTCGGATCCTGACAAACCGCC GGACCATGATGCGGTTTGTGATGCCTTCAAGCAGCATACGTGTGTCGACTAT[G/A]CTCTTC ATTTTTATGTTCTGGCTTCCCTACACTTTGGATCCAACGCTCAACTCATACCACAAGATGA AGGTACTTAATTCATTTCCATTTTGTTTTTTTTTGTTATGATTTTGTATGTTACTATCTCT GTTCGAAACACTTTTGTAACTATATATAAAGTGAAGTGTCTCTTAA
>Gm13_3 5815670	GATGTCCGAGGGTTCGGCCACGCGAACTGCGGAGGGCTCATATGCGTATGAATCAAACCG CTGAAACCCCTGGGCAAGCCGGTCAGCCAGGTCCTGCCGTGACGCTGCACCTCTTCCCCT GCCTCAAACGACCTTTCCACAGACATTTTCTTCTTCTTCAACAAATTCATAACT[T/G]C CCATAAGACCTAGGGTTTCACTTATACTCCCAAATCTAATTAGAAGACAACAAATAAAAGCG TAAACGAAGTGAATCTAAATCTAACTCTTTCTTATCTAATCAAAGGAAAGAACAAATTCGTCT TATTCACAAGGGTTTAGGGTTTCGGTTTCAGTAGTGTGCAAATCTAATTTAA
>Gm13_3 5819710	ATTCCTATTGGTTACTAGTAATAAAACCTTTACACTATCGACTATCAATCCACGTTCAAATT GATCATGTAATAATGGTGGAGTACCTGGTCTTTGAGTTACCAAGTGTGAGTTCAAGATCATC TGATCCAAATTCCTCATGGATCCTTTCTCCTTCCCAAGGCTTGACAAGCCCCAA[C/T]GTGTT GCTTCCAAATGCAAATTCATCTGAAACAGCTTCAGACATAGGAATGTCAGCATTTTGTATCAG AGCCAGGTGGAATAGCAGGAGAGCATGTTCCACTCTGAGCAGGAGTCCACATTCGCGAAC CGCTTCCGGGCAAGCCATCTTCTTGAAGGCAAAAGGGTTTGAGGAGACCAAG
>Gm13_3 5823484	AGCCTCGTCCCCGACGTCATCGATCTCAGACCTCAAAAAAACTCAAAAGGGTCCTCCTCAA ACCTCCCAAGAACACAACTTTGGACCAGATTTCGATCCAACAAAGTTAAATAGCCTTCTGG GTCGGTCACAAATTGAAAAGGGTGTGTGTGTGTTAGAAGAAGAAGGTTGTGTTATAA[A/G]A AGAATGTGGAGGAGATGGTTGGAAGTGGCAAAGTCAGCACATTTGGTGGATTTGGGAAGT GGGTTTGTAGCCGGAGAGTGTGCGGAGATCGGAGGAGATCTGGGTCCGAGAAATGGGAA ATGGTTTTGATGAGAGTGTGTGTGATGTGTGTGAGAAAGAAGCGGAGGAGAAGAAGAAG
>Gm13_3 5823512	GACCTCAAAAAAACTCAAAAGGGTCCTCCTCAAACCTCCCAAGAACACAACTTTGGACCA GATTTCGATCCAACAAAGTTAAATAGCCTTCTGGGTCCGTACAAATTGAAAAGGGTGTGTG TGTGTTAGAAGAAGAAGGTTGTGTTATAAGAAGAATGTGGAGGAGATGGTTGGAAGT[C/T]G CAAAGTCAGCACATTTGGTGGATTTGGGAAGTGGGTTTGTAGCCGGAGAGTGTATCGGAGA TCGGAGGAGATCTGGGTCCGAGAAATGGGAATGGTTTTGATGAGAGTGTGTGTGGATGT

	GTGTGAGAAAGAAGCGGAGGAGAAGAAGAAGAAAAGGAAAAGAGTGAGGAGGAGGAGGC
>Gm13_3 5823533	GGTCCTCCTCAAACCTCCCAAAGAACACAACCTTTGGACCAGATTTCGATCCAACAAAGTTAAA TAGCCTTCTGGGTCGGTCACAAATTGAAAAGGGTGTGTGTGTGTTAGAAGAAGAAGGTTGT GTTATAAGAAGAATGTGGAGGAGATGGTTGGAAGTGGCAAAGTCAGCACATTTGGT[G/A]G ATTTGGGAAGTGGGTTTGTAGCCGGAGAGTGATCGGAGATCGGAGGAGATCTGGGTCGGA GAAATGGGAAATGGTTTTGATGAGAGTGTGTGTGGATGTGTGTGAGAAAGAAGCGGAGGA GAAGAAGAAGAAAAGGAAAAGAGTGAGGAGGAGGAGGCACCTTGATTAGCCAAATTTGCG
>Gm13_3 5823553	AAGAACACAACCTTTGGACCAGATTTCGATCCAACAAAGTTAAATAGCCTTCTGGGTCGGTCA CAAATTGAAAAGGGTGTGTGTGTGTTAGAAGAAGAAGGTTGTGTTATAAGAAGAATGTGGA GGAGATGGTTGGAAGTGGCAAAGTCAGCACATTTGGTGGATTTGGGAAGTGGGTTT[G/C] AGCCGGAGAGTGATCGGAGATCGGAGGAGATCTGGGTCGGAGAAATGGGAAATGGTTTTG ATGAGAGTGTGTGTGGATGTGTGTGAGAAAGAAGCGGAGGAGAAGAAGAAGAAAAGGAAA AGAGTGAGGAGGAGGAGGCACCTTGATTAGCCAAATTTGCGGATCAAATTCATGATCACGG
>Gm13_3 5835277	TCCTCTTCTCGCCGAAAAATCAAAGAAGCTCCGAAAAGAACAAAGGAACAAAAAAGAAA CACACACACGCGAACACGAATGTTAGAAAGAAACACAGCACACAGCAGTAAGCGAAT TATAGAGGAAACGACATCCTTTCTGCTTTTTATATAAGGTGAGCTAAAACGACGAC[G/T]T CGGATTTGAGAGATTTGGGATTGTGATCTGAGTAGATCTAGGCGAGGGAGGTTGGGTGGT GGAGGCGGAGGAGCATGGTGGCCGAAAAAGCCGCCGAAAAAGAGAGGAAGTGGGTTTGT GTTAGGTGTGAACGTGCACAATGGGGAGAAGAGAAGGTTGAAGGAGCGAAAGAGAAAGA
>Gm13_3 5849515	TTTTGTGCTGTCTTTGAAGGTCTTAATCTCTTGCTCTAACTGCTTCAGGTTGTTCTTGAGTTC GGTTAGGGTTTCGGAGGCAGTTTCTTCTTCTTCTTCTTCTTCTCGTCGGAGGGGGTGGGT TTTTGGGAGAGGCCGAGGGCTTGGGTGAAGAGGACGTGGAGGGGCTTTTTGGTGG[G/A]C CCGGCAAGGGTTGGGGTTGGGGTTGGGGTAGGGGCGGGCTGGGAGGAGGTGGGTCTGG AAGGGATTTTGAAGAGGGAAGGGAGAGGGTGCGGCGAGAGAAGAGGGTACTGAAGATA CCCATGGAGGGCGCAGTGAGAAACCTAGAACGAAAAAGGGAGGCTTTATTTATGGGCTTC TA
>Gm13_3 5862124	ACTAATTAATGTCTTATGCTACCAAAGCACTCATACAATCACTAACAATGACGAACACTCAA AATCACTAGTCTTAATTAATAACATCTATCAAATATGTTTTCTGAACTTTGCAATTTCTAATTC TCTAATTGCCCAATTTCTTTGCCGCGTTAGCTTCAACCCTAGCCTCGTCATTT[A/T]TTTTGCT TCCGTTGTTACCATTCGGCATTTTTTGAACCACCTCATTATGTTGTCCACCTTTCCCCTGGC TGATGGGTGACCCTCTTTCCCTCCAACGCTTCCATGGCTGGGGCCTGACCCTCTTGTTCC TCCGTGCGTTCTCACTCTAAACAACAATAACGTCAGTTCTTAATTCATAT
>Gm13_3 5862205	AACATCTATCAAATATGTTTTCTGAACTTTGCAATTTCTAATTCTCTAATTGCCCAATTTCTT TGCCGCGTTAGCTTCAACCCTAGCCTCGTCATTTCTTTGCTTCCGTTGTTACCATTCGGCA TTTTTTGAACCACCTCATTATGTTGTCCACCTTTCCCCTGGCTGATGGGTGAC[G/A]CTCTTT TCCCTCCAACGCTTCCATGGCTGGGGCCTGACCCTCTTGTTCCCTCCGTGCGTTCTCACTCT AAACAACAATAACGTCAGTTCTTAATTCATATTTATATATATATGACAGAACATCTAG CTAAGGCTGAATGAGATTTATGCATGTAAAGAGTGTTAACTTGATCATC
>Gm13_3 7094557	AAGACAGCCTTGAAACCCTGTCTTCCAAAGGAGCAGGTTAACATAGCCAGTGCGCAAT GAGTTGGCAGAAGAAAGGAATGCTTGATCAAATTATGGATCAGAATCTGGTGGGGAAAGTC AATCCTGCTTCTCTTAAGAAGTTTGGGGAGACAGCTGAGAAGTGCTTGGCCGAGTATG[C/T] AGTGGATCGGCCGTCTATGGGCGACGCTTGTGGAATCTTGAATATGCTTTGCAGCTTCAG GAGACCTCATCAGCACTCATGGAACCTGAAGATAACAGCACAAACCACATTACTGGAATTC AGTTAACACGTCTCAAGCCTTTTGATAACAGTGTCATATGGTTGATGGGGGAAACTC
>Gm13_4 2335929	GTCATGGTGATCTAAGATGAACAACTTCTTGTTTTGGATAGCCTGTTCAACCAAGTACAATT AATTAATTGCTATTTCATTGGAATTACATTAACAAAAACACACACAAATAGATAAAAAGCAA TAGAAAAACATTATGGTGAAACAATTTTAATGTGAAACCAATACTCATATCA[T/C]TACCTG CTCAACAGTGAGCCACCTAAGTTAGGCTCCAAATGTTCTTTTGCTATTATACAGGTATGAT CACCATAGGCTTGAGTATCTAGCTGTCTAGTACTAGTGGGAACCTCTGGAATAAATTTATCAGC CAATGAATGAGGATAGCACTCCCAAATTGACTCCAAGCTTTTGTTGCGT
>Gm14_2 683857	CTTGTTTCAATGTTTTATTTATTTTTATTTTTATTGTTGTTCTTTATTGCCTTAAATTTTA AGAACATATAAGCCTTATTCATTCTCATGTTTCAGGTATGTCATGAGCCCCCTATTAGGAA ACGAGGACATGATAAGGCCCTTCAAGCTGCCCTTTCAACCGCGTTTTGCA[T/C]GTTTCAC ATTATGATTGAGGCTTAGGTTCAATAACCAATTAAGATGAGGGTTTGAGTTTCTCAGGATA GTCTCAGCTATGAAATACACCCTTCAAGCTTCCCTCAGCTCAATTTTATTCGTTTACTTAGC TTGTTTATTTGACCAAAGGATACCTTTTTGAAATTTGCAGCTGGTAGG
>Gm14_2	TGTTTCAATGTTTTATTTATTTTTATTTTTATTGTTGTTCTTTATTGCCTTAAATTTAAG

683859	AACATATAAGCCTTATTCATTCTCATGTTTCAGGTATGTCATGAGCCCCCTATTAGGAAAC GAGGACATGATAAGGCCCTTCAAGCTGCCCTTTCAACCGCGTTTTGCAGG[T/C]TTCACAT TATGATTGAGGCTTAGGTTCAATAACCAATTAAGATGAGGGTTGAGTTTCTCAGGATAGT CTCAGCTATGAAATACACCCTTCAAGCTTCCCTCAGCTCAATTTTATTCGTTTTACTTAGCTT GTTTATTTGACCAAAGGATACCTTTTTGAAATTTGCAGCTGGTAGGAA
>Gm14_3 559049	AAGTCCGAATGTCCCAAAGAGCTTGAGCCAGCAGATAGGAGGACTGCTGTCCATGTCAAC CTCACAAGGCGGGATGAAGACTACCCTGTATTCCCTTTTCTATTTTCTAGTTTTTGCATCTA TCTATCATCCTATTTAAGGTTTTTACATTTTCTTTTTCTTTTTGCCAAAATGTATG[G/C]AGGA ACCAGTGAAGCCACGCCATCGTGCTTTCCAGGGTGTGGAAGAAGCTTTAGGTAGCACCAG CTCCAGCAATGATGAACCTATTCAAACAAGTGGTGCTTCACCAAGTACTGCTCCATTGCCAA CAATGGGTTTAATTGTAGATGAGGCACAGTCAGTGACATCAATCCAGCTGAGGT
>Gm14_3 559050	AGTCCGAATGTCCCAAAGAGCTTGAGCCAGCAGATAGGAGGACTGCTGTCCATGTCAACCT CACAAGGCGGGATGAAGACTACCCTGTATTCCCTTTTCTATTTTCTAGTTTTTGCATCTATC TATCATCCTATTTAAGGTTTTTACATTTTCTTTTTCTTTTTGCCAAAATGTATGC[T/G]GGAAC CAGTGAAGCCACGCCATCGTGCTTTCCAGGGTGTGGAAGAAGCTTTAGGTAGCACCAGCTC CAGCAATGATGAACCTATTCAAACAAGTGGTGCTTCACCAAGTACTGCTCCATTGCCAACAA TGGGTTTAATTGTAGATGAGGCACAGTCAGTGACATCAATCCAGCTGAGGT
>Gm14_2 7341132	GGTGAAGTCAAGATATGAGGCTGTATAGAAGCGGATGCGTGACGTGGAATGGTGATTTGT AGATATGAGGTTGTATTGTTACGTCTTGCTTGCTCGGGAATTTGAAGCCATAGTTTGAAAAT CCAACACCTTGAGGACAAGGTGTTTTGATGGGCTCGGGAACCAATGTTACGTCTT[A/C]JAA ATTCGAGGACCAAGACAAGCTTATTGAGAGAGAAGAAGAGAGAGCTTTTTTGATATTTCTTT CTTATACATATATATTGCTAGCTGCTAGGGAACAACTCCCAACAACAATAACAAAAGGA TAAGCGCAATCATGGAATATCCTAATAGAATGATATCCATAACATTCTACACTA
>Gm15_2 144200	ATTCACCTCATATGTAACCTCTGGGTTTACAATTTCAAACACTACAAGACTGATCGCATTTTC TTTGCCAACAATGTAAGCTACTTAGTGCTTACCATGAGAACCATAAGACACCCTGTAT CAAAAATTGAAGTTATTAATTATGTAATGCAGACATATCTTCCAAGCGAGACAC[T/C]GGCTC CACTTGTCAGTACAGAGAAGAAGAATTGAAGAATGTAAGAGGGGATGGAAGTGGTGAGC GCAAGGAATGGGATAGGATCTATGATTATGATGTCTACAATGACTTGGGCGATCCAGATAA GGGTGAAAAGTATGCACGCCCGTCTTGGAGGTTCTGCCTTACCTTACCCTCG
>Gm15_1 2301810	TGACATAATATCTAAATTAATAAATTGGTTTCTATGCTATTGTCTTGATTCTTGAGTCTTTT GTTCTAATGAGGTCTTACCTACATAATTTTCAAGAGGCCTTTTAAATTGTAATACTTATGATAT AGTTTTAGTTGAGTGAGTTGAACCTTGATACATTTTCTGTAGGTCATCAGCG[T/C]CCACTTA GGATGGGTGAGATCTGTTGCAGTTGATCCCAGTAACACATGGTTTTGTACTGGTTCTGCAG ATCGAACTATCAAGGTATTTGATCACAATAATGCATGCCATAGTTGCTTGTATGAGAAGGCA TGTTTATAAGGAAAATTTGGTGTCATGTTGAATTGTTGATTCTATGGGT
>Gm15_1 3886434	TACAATATAATTTTTTAAACAAGAATAGTTTAAATAGTAATAATCATTTTTTAAATAAAT TTGTGTAATTGGTAGTAACAATTTACTTACATTACTTTAAACAACAATAATGTGAAACAGAGG GTTACATGCCATCAGGGGCAGACGCCCCCCCCCCCCCCCCCCCCCCCCCCCC[G/C]CCCAC ATCACAATTCACAAGTCTCTTAATATTTTCAAGTTAAATAAAAATTGCCGCCCTCTAATATTTA TGGCTTACTCTTTGTATTATGATTATACACCATTATATCCAATAATTCCCATCCAAATTGATAT TAAAGTTTGTCACATGGCCAAATGGACCAATCCTGTTCTGTTACTTCAA
>Gm15_4 8317889	ATTCACGAGTTCGCGAGTTTTCTGTTGTGGGACAAAAATAAACTATTTTTTTATTTTTATTCC CGGGAAACAAAACAGAACAGAAAATACCCAGAAAAAAGAAAAGAAAATTGAACGAACCTGA AGGTGCGAGTCTCTTCTCCCAGCCACTAGGAACCTGCAAATCGAGGTCCAATTCTT[T/G]GGA TTCAATTGAGGAGGACCCACCAAGCAAGTCCCAGTAATCAGAGCCGTTGATTTCTACCCG TGTGATTCAATCCCCACCGTCCGATGCTGCTCCTCTGTTAGCCACCACCCATCACCCGAA TCAGAGAGCTAACCTCAGCAGCCATTGCTGTGCGCGTTCTCACAGAGGATTCTAG
>Gm15_4 8317898	ATTCGCGAGTTTTCTGTTGTGGGACAAAAATAAACTATTTTTTTATTTTTATTCCCGGAAACA AAACAGAACAGAAAATACCCAGAAAAAAGAAAAGAAAATTGAACGAACCTGAAGGTGAGT CTCTTCTCCCAGCCACTAGGAACCTGCAAATCGAGGTCCAATTCTGAGATTCAA[A/C]TGA GGAGGACCCACCAAGCAAGTCCCGAGTAATCAGAGCCGTTGATTTCTACCGTGTGATTCA TTCCCCACCGTCCGATGCTGCTCCTCTGTTAGCCACCACCCATCACCCGAATCAGAGAGC TAACCTCAGCAGCCATTGCTGTGCGCGTTCTCACAGAGGATTCTAGAGAGAGAGA
>Gm15_4 8317931	CTATTTTTTTATTTTTATTCCCGGGAAACAAAACAGAACAGAAAATACCCAGAAAAAAGAAAA GAAAATTGAACGAACCTGAAGGTGAGTCTCTTCTCCCAGCCACTAGGAACCTGCAAATCG AGGTCCAATTCTGAGGATTCAATTGAGGAGGACCCACCAAGCAAGTCCCAGTAA[G/T]CA GAGCCGTTGATTTCTACCGTGTGATTCAATCCCCACCGTCCGATGCTGCTCCTCCTTGTA



	GCCACCACCCATCACCCGAATCAGAGAGCTAACCTCAGCAGCCATTGCTGTGCGCGTTCT CACAGAGGATTCTAGAGAGAGAGAGAGAGAAAGAAGATAGAAAAAATATGGAGAGAGAG
>Gm16_1 37436	ATAGGTTCTTAGGAAGCTTGTGGATGATGCTTCCAAGTTTTGAACGATAAGGTTACCAAGG CTGTTGTTACTGTGCTGCTTACTTCAATGACTCCCAAAGGACTGCCACCAAGGATGCCGG TCGGATTGCTGGTCTTGAGGTTCTTCGTATTATCAATGAACCAACCGCTGCATCCT[T/C]GG CCTATGGCTTTGAAAAGAAAAACAATGAAACAATCCTTGTTTTTGACCTTGGAGGCGGCACC TTTGATGTCTCTGTGCTCGAGGTTGGTGATGGAGTGTTTGAGGTCCTCTCTACTTCTGGTG ACACCCACTTGGGTGGTGATGACTTTGATAAGGTACCTTTCCCTTCTTTCTTCT
>Gm16_4 318195	TCAGCGATCAAGGAAACCAGACCTCGAAAAATTATACTTCAATCAAGAGTTGAATAAGCCAA AAACAAACTATACATATATCTTTTAAAAACGCATATATTGTTATATATTACAATGATAAGATA AAGATTAACAGAAAGTCAATTGGTCCCGTTTAAATAGCTGAAATGTTGCAGACT[T/C]JACCTCT TATGCTCTCTGATAAAAGGTTGGCTTTCCACGTCTATCCCAACCGATTCTTCACCAGAGCCA GAGCTGTTTACAAACAAACAAATGATAAAGGAGAGCTCGAATCAGGAGGGAAAAAAACAAA AAATAGGCTCCAAAGGTTAAGAAAAAGGGTCTGCAACTGCAATTGCGGCC
>Gm16_1 9670636	GAAAATACTATCTGGGTCTAGGATCAAAAATACCAACATATCCCTGTAAAGGGGGAATTTGGA ATACTGGAACCTATTGCAAGAGGCGGCAACAAAAAGTATCAAGCTCTACCAACAACCTTGAA TAACAAAAATAAGTGTAATATGTAAAAACTCATGTAATGAGGAGGAACGAAACGG[G/C]ACC AATCCTTCTTGCCAGCCCATGTTGCGAAGCATCCGATTACCAACATTGTTTTCATCAATTGC TTTGTCTGCCGTAATCACCTCAAAGTATCAAGGTTGGCATCTCCAACATATGCGTCCTCCAC CAACCCAGGAGGGAATGGCATGGGATCACCTTCTTGATGCAAAATCTCGA
>Gm16_3 0464934	TCTCTCCCTCCTCATGAGGAGAAAAAAGAAGCTTTCACTTTCTTATTATTCATCCAACACC ATCTCAAATCGAAAGGAACAAGCTTCACCATTCACTTCTTCTACCTCCTCATGTAGTA CCAACCAAAACACCACCTTCTTCCCCCTCCCATGAGGCAAAATAACTACTTCTGA[T/G]TTTG TTAATAGCCAGAACAATCCATCCCAGGCTACAAATCAAGGTAACTATATATTATTGCCAGG TTGATTATATATCATATTATTCTAATAATTCAATTACCATCGATGTTATTATATGGTCCTCTC AATTATATATATATCATATCATATCATATCAATAATCTGGATCCACATCT
>Gm16_3 0493681	GAACCTTCAGTTCATACCAAAGCCAGGATCGAAGTGCTTACTTCAATCACTCTTCAAAGGA GAGGGATTGTTACCTTATCATCATGATCAGAAAGAAGCTGAGTTAGATTTCAAACTTCAA CAATGTTATGATGGGGGGTGGCCAATTTTCCAGTCATTTCAAGGAGCCATTGCAA[A/C]CGT CGCTGACATTGGACCAGGGGGCGAAGGAGAGCTACTGAGGTTTACATGCCAGAAAAACATGT CAGAGAATATTTATTCGGATGAAGGAAGGTATTTAATCCCCAGACAAGATCCATTGATCCCC AGGCAAGATCCATTGATCCCCACGCAAGATCCATTGATCCCCAGGCAAGATCCCT
>Gm16_3 0493887	AGGAGAGCTACTGAGGTTTACATGCCAGAAAACATGTGAGAGAATATTTATTCGGATGAAG GAAGGTATTTAATCCCCAGACAAGATCCATTGATCCCCAGGCAAGATCCATTGATCCCCAC GCAAGATCCATTGATCCCCAGGCAAGATCCTTTGATCCCCAGGCAAGATCCATTGGT[G/T]C CCAGGCAAGATCCTTTGGCTGCAGTAAATATGACTGATTGGGCTGCTAATAATGCTCGCAT AGCAGGGCCTTCCCAATCTCACTTAAATACTGGAGATTTTATTGGTCATCATTGGTTTCCTG CTGATCATCAAGTTCGTGGTGGTGGCTGGAATGGATCTGATGGTGGTGGTCTTTCA
>Gm16_3 0517229	GAGAAATCAGCAAAACCATTTAATCAAACCCCCAAAAAGAAAGGCAAAAACAATAAGTTGC TACCAAAAAAATATATATGAGAGAAAATCCTCCGAAAAATTCTTTTTTTTTTATGAAATTTTG GTTGCATCGACAACATAAGGGGGGGGAATGCTCACAGTGGCTGCCGGAGACGGC[A/G]GCG TGCGGTGGCTGAGAGTGAGAGGGGTGAGCTAAAAGGGGTTGGTGAGGCTCGGAAACAAA GAGAGGAAATTAAACGATCGTACGATGCACCACTAATTAGGAAATTCGTTTAAATCGAATCA AATCAAATTGATTTTACAAATCTACTTTAAAATTTAAATTTTAATTATAATAAAAA
>Gm16_3 0517232	AAATCAGCAAAACCATTTAATCAAACCCCCAAAAAGAAAGGCAAAAACAATAAGTTGCTAC CCAAAAAATATATATGAGAGAAAATCCTCCGAAAAATTCTTTTTTTTTTATGAAATTTTGGTT GCATCGACAACATAAGGGGGGGGAATGCTCACAGTGGCTGCCGGAGACGGCGGC[G/A]TGC GGTGGCTGAGAGTGAGAGGGGTGAGCTAAAAGGGGTTGGTGAGGTCTGCGAAACAAAGAG AGGAAATTAACGATCGTACGATGCACCACTAATTAGGAAATTCGTTTAAATCGAATCAAAT CAAATTGATTTTACAAATCTACTTTAAAATTTAAATTTTAATTATAATAAAAAAT
>Gm16_3 0539483	GGTTTGCGGAAACTGCCAGACTGTGTTGTGCCAACCAACAGGCGGACGGGCAAGGTTAAC CGAAGGGTGCTCTTTTAGGAAGAAGGGAGATGAATGGTTGTTAACCGACTTTTCTTTGT CAATGATTGAAAATTTTCGGTTGTTAGGTTTTGAAATGGAATCCATTTCGTTATG[G/A]TAT ATTAATTTTGTAAAATTTGTAAGACAATATCTCTCTCAGCCTTCCACAAGACATAGTAGCCAAT GTTGAATTTTCTTTATTAAGTGGATGTGGTTTTGTTGACAACATTTACTAAAATCAATAT GGTTTAGTTTGATGGTTATTTTACAATAACATGTTTGTATGCTCTGATG
>Gm16_3	CAACAGGCGGACGGGCAAGGTTAACCGAAGGGTGCTCTTTTAGGAAGAAGGGAGATTGAA

0539518	TGGTTTGTAAACCAGACTTTTCTTTGTCAATGATTGAAAATTTTCGGTTGTTAGGTTTTTGAA ATGGAATCCATTTCCGTTATGGTATATTAATTTTGTTAAATTGTAAGACAATATCT[T/C]TCTC AGCCTTCCACAAGACATAGTAGCCAATGTTGAATTTTTCTTTATTAAAGTGGATGTGGTTTTG TTCGACAACATTTACTAAAATCAATATGGTTTAGTTTGATGGTTATTTTTACAATAACATGTTT TGTATGCTCTGATGACGTGTTAGTGTCAATAATGTACATATGAATTTTG
>Gm16_3 0539519	AACAGGCGGACGGGCAAGGTTAACCGAAGGGTGCTCTTTAGGAAGAAGGGAGATTGAAT GGTTTGTAAACCAGACTTTTCTTTGTCAATGATTGAAAATTTTCGGTTGTTAGGTTTTTGAAA TGGAATCCATTTCCGTTATGGTATATTAATTTTGTTAAATTGTAAGACAATATCTC[C/T]CTCA GCCTTCCACAAGACATAGTAGCCAATGTTGAATTTTTCTTTATTAAAGTGGATGTGGTTTTGT TCGACAACATTTACTAAAATCAATATGGTTTAGTTTGATGGTTATTTTTACAATAACATGTTTT GTATGCTCTGATGACGTGTTAGTGTCAATAATGTACATATGAATTTTG
>Gm16_3 0552971	CTGGATGTGCAAGGCATTGGAGGTGTTTTAAGAGTTATGGCAGAGGAAAGAAAGCTAAAG AAGGAGGAAAAATAATGTTGATTCTTCTTCAAGTTCAGTGCTAGATGTGTCAGAAAATCAAA TATGTCATCCAAATACTATAAAAGAACAACCTACCACACAACACCATAGTATGTCAT[G/A]AAG AACCAATATCCTGGCCCATGCTTCCCAGGAGATAGCTATCCATTTTGTTATTGACACTACT CTTGCAAGTGTGCTTCTTTATGTGGCACAGTAACTTCCATTTTGAAAAAAATATATATTTGTA ATCACACTGTCACAGTTGGTTTGCTTGAACACTTCCAACTTAATTTAGGGT
>Gm16_3 0676914	TCGGTACTTGGACCTCATCTCCTCCATCAGGGCCCGAGTAGTGGGCCCGGTAGCACCGCCC CAGCTGCCTCGTTCGCCGCAGCCGCCGCCGAGCACCTCCTCCCGCGTCAGCACCCGCG ACTTCGCCAGCGCCGTGTCGCCCTCGCCGCCCTCGATCGCCACCGGCCACGGCGGAGAG G[G/A]TGCAGAACCCCACTCGTCCATCGGTGCCGAGTCGCCAACTCGGAACCGGGTTTCTT GCTGGAACCTCGGAAGAGTTCGTTACGGTAATTTTATTTTCTGACTTTGATTCTTCGGCGTTG GGAAACATTTAGAACCGTTGGATACCGTTTAAAAATGATCTGACGATCAGTTTGAAGTAAT
>Gm16_3 1154583	TTTTCTGGTTTGTATAAGTTATTCTAGCTGATTTTCTCACCTCAATTGAATCTTGATACCAT GTAGAAAAATAATAGCAAGAGAGAGAAAAAACAGAATACAAAGCTTCTATGGTGAAT GAAAGTGATTTTCGTTCTGAATTTTCCAACATACATACAGTGACTTGATTTTTT[A/T]ATTTTT TTTGTGAAAATTCAATATCACAAGCAAAAAGTGTGACATTGGCCTAGCATCCCTCAAAACAT GGGTTACCACACTAGTTCACCAACAAGTCGACTTATTACATATCACAAAATATTAACAAG ATATCTAATAGATTTTACTTCCAATATATCAAAAAAAGCTCATCAATTC
>Gm16_3 1154850	CCAACAAGTCGACTTATTACATATCACAAAATATTAACAAGATATCTAATAGATTTTACTT CCAATATATCAAAAAAAGCTCATCAATTCTAACAAGGAAATACAAAGGAAAGTGCTTCGAA GTACTAAAGGGTCTTTACTATTTACCAAGGCTTCCTTTTCTTGAAAGAACCTT[A/G]TATCT GCAGAGGATCAAAGCACAACAGTTCATATTATTATTATTGTTGCAATATCAACTTTTT TTGCCTCCAATTATCTTTTTTTTCCATTTCCACTAACCCCCCATGAAGAGGAAATTGACTCA AATAGTGGGGTTGCTCCTGAATGATCCAAGAGCCTTAGCAGCACTTGCT
>Gm16_3 1189225	ACTCCTTTCTGGGTCCCCGACATGGGCGCTACTACTAATAACGGATTTTCAACCATTTCA CACTCTCATAACTCATTCTTCAAAACAACAAACCTTAAAAACAACAACTACTAAACCTTCTC CTTTTCTTTTCTTCTCCATTTTCCCAACTCACACTTCTCATTTTCTCAGGGAAAT[C/A]GAAAA AGGAAATGATCGGAACTTCATCGACGACATTGACTGCGGCAACTTCTTCGACCACATCG ACGACCTCCTCGAATTCCCCGACGACGCCGCCGCGCTGACACCTCCGCCGCCGCTCCTG TCCCTCCACCGGCGAACTTCTGGTCCGCCGAGTCCGACTCGCTCCCCGCCACCG
>Gm16_3 1191863	CAACCCCAATCCAGTGATGCAGCTCATCTCCCCTGCATCCTCCACTGGTGAGAACACGCAG CACAACGCTGCCAACACCTCCAAGGCATCATCGGATTCCGAGAATTCGCTGAGTCGGTGA TCAAGGCTCCTAAGCAGGCCTCTGGGGAGCACAAGAAGAAAAAGAAGATCAAAGTGA[T/C] GTTCCCATCAGGTCAAGAGCGGAATGCACCATCACAGGCAATTAGGAAATGCTTGCACTGT GAGATAACCAAGACACCACAGTGGAGGGCAGGGCCAATGGGGCCGAAAACACTCTGCAAT GCTTGCGCGTGCGCTACAAGTCAGGCGGCTTTTCCCGAATATCGCCCTGCAGCGAG
>Gm16_3 1204451	CCCAATCGAAGCCCCACCCATGACTGAGCATGCAATTGCAATTGCAATGCCAGACACACT CTGAAGGATACGCAGGGCATGCCCGGCACCCTCGGTGGCTTCTCTTGCGCTTCGCCAG TTCTCTTTCGCCCTCGTTTCCCTCTCCGTCATGGCTACCACCTCCGATTTCCCTTCTG[G/A]C ACTGCCTTCCGGTATTCATTAATCCCCCTTTCTTATTTTCTGTTAATCGTGCTTATTTTT AATTGGGTTTGTGGTAATTGTTGATTTGGGCTTGTTTTTGTGTTCAAAATCAGCACCCAG ATTTTATTTTTTTTCAAGTGGGTTTTATCAAGGAGCAATGAAATGAACCTCTT
>Gm16_3 1225684	TAACAAAACAGATCATATATCTGCCATTATTCTAAAACAGAAAAGGATAGTTAAAAATAAAGAA TACAAGGTTACTCCAGAATAACGATGTAGTAAAAGGAAAGTATAATGTACAAAGGTTTCTAA ACAATACAATATGGTGAAACAAATAGGATACCTGCATAATTTCAAGGTGCATCAG[T/G]CCCA CACATTGAGGCTCCGAGAACCTTATCTGTCTCAGCATCAACAACAAGCTTCATAACAGTTTT

	TTCTTGTGCGCTATTTGAAGAATATCAGCAACCATAATCTAGATAAGAAAGTTATGCTGCAA AGAACATAACTAATACTGGGTCCAACCTGAGATTGAAGAGCATGTGATGATAA
>Gm16_3 1229414	ACAAAGTGATTGATTTTGTATTTATCATTGTACCCAACCTACCTATTATAAATAGCATTCATGT TTGTGCCTCCAAGACACAACGTTTCAGATATTTCTTCTCTATAGTATCCTTACCAGTGGCAAA CAGTACAGCATCCGCAATCAGCTCTTCACCATGATCTGTAATAACTTTAATGC[C/T]GTCCTC TGTTTTGATCAACTAAAATAATGAAATTTGTTACCATTAGCTAAGCCTCCAACCAAAGCAAAT GAAAAAGTTCTACAAATGTGAGGTTGCAAACCCCTAAACCCAAAGGAAATACCTGTGTCAAA TTGGTCCTTGGATGCAAATTAATTCCCCTGCCTTCAAGATTTCTTGCCA
>Gm16_3 1232875	CATACAAATCTAACATCTTATTGTTTCATAGTCTCTATTTGTTATTTTTCTCCTACCATAATCCT AAACTTGTAATTCTTATTAGCATGTACATATCAAATATTATACTCACAAACAGAAACAATCAT GTAAGTAAGATCATTTTGCAATGATATTTCAAACAGGCAAATTCACCTGTC[A/G]AGGAATA TTTGGACGTTGAGCCCTGCTGCCAGTTGCAATCAATATGTGCTTTGCTGTGTAGGACAATTT AGTGCCATCCAATTGTGTACCTCAACTCATTTGGACCACTATCTTTCCCTCGCCTTCAA ATAATTTAACCCCAATTGGATAACATCCGCTTGTAAGTCCATTAA
>Gm16_3 1292709	GGCACATTGCTCTTTTCCAAGGGCATGCACCAAGCCATTCTTTGGATGACACTAAAGGGGC AGGGGTAGCTGAGGCACCCATTTCATGATAGCAAAGAGAACAACAGTAAGGTGCAGAGCAT GAACCTCACTTTTGGATCAAATCCTTCAAATGCAAATGAAGTTCCAAATGCCATGGAT[C/T]T AAAAGGCGATAGCGATGATCCTATCAATTATTTCTCATCATTGAACCTTTGAAGCCAACCTTG GTGAGTTTAATGAATTCACCAGACCTCTATTTGATGACCCAGAAGGTTCTGTTCTTTTGTG ATGCCTGGATCTTTCAAATCTGGAGGAGAATCTAGTTCTGACAATCTTTTATAAC
>Gm16_3 1363451	TCGGGTGCTTTTCGATATGTCGTCGCCCGCCGCGTATCCTATATCGGCTGCGTAACCAATG AGGAACACGGCGACGGCGACGGCTAAGGCGCCCCGAGAATAAAGGGGCGCCTGCGACC AAAGCGGGAGGTACAGTGGTCGCTGTAGTAGCCACAATGGGCTGTACCACGAGCCCGG[ G/A]AATGGGCCCCGAGAGCCAGATGAAGGAGGCCGCGCGTGGGGGACTCCTAAAGCT GAACGTAGGGGGTTAGAAGAGAGAGCTGGAGAGCCCAACCGAAGTGTATGCCGGCGGCG ATGGACGCCACGGCAAACATTTTCGGAGTGGGCTCGCTTGGGCCGGGCCGCTCCAGA CTAAG
>Gm16_3 1427024	AGATTTGGATTGGAATGGTTTCCCCAAGCAAATAAAATTAAATTAAAAATAAAAAAGACAAG CCCTCCTCAAAATTATCATAAAGCAGTTAACTCACAAGCCTCAACCCCATTCATCCTACCC AACAAAGAAAATCCCAATGTCATAATTGGTGTCAATTCATTTCTCTAGTGTGTGG[G/C]ATCT TTTTCTTAGCATGTGTCATCTTTGTTGTTCCCATTTCCATTACCAAGAGCCCTTTGCTTTC TTCATCTCTCTCATTTTCCTTATTCTAAGCCCATTCGAACTACCCATTTCAATCCTAGTGAG CTTCTCAGCCTTCAAAGCCTCAGCTTCCCAATCTGTCCTCACCACACAA
>Gm16_3 1427304	ATTGCAACTACCCATTTCAATCCTAGTGAGCTTCTCAGCCTTCAAAGCCTCAGCTTCCCAAT CTGTCCTCACCAACACAACATAGAGGATTGACACAGCACATGCCACCTGGGCCGACAGAA GCCCAAACCAAAGCCCACTGAACCCACCTTGAACCAAAAAGGCCAGGCCACGGCCA[G/T] TGGGGTGCCCAAAAGTAGAATGAGCCTAGGTTTATGTGGGCCCAATACCAGGTCGGGC CGTGCCACGAAGGATCCACAGCCCGTGGTTTGTGGACAGTTCCCAAGCTCACACAGGCC CATAATCGGCATGACTGATGCAACCAAGGCTTTTACGGGCTCGTCGTTGGTGAAAAGCCC
>Gm16_3 1459969	GAATCAACATGGTGTGTATTATGATTGATTGATAACCTTGCCCTAAGTATGTGGTGAGAG GCCTGATCCCGAGGATTTTGTATTTAGAAAAATATCTGTTGAGAGAGGTGAACCCCTAAATG AATCTTAGTACTTGAGGGATTAGCCCTGGCTCCGAACGGGAAGATACCTGAGTGC[G/A]AC CAAAAAAAGAAAAAGGAAGAGTAAAGTGTGGGAAGAACAAAAAGAGGAGAAAAATAGATT GAGGGCATCTTGGTTTGTAGAACTTAACTGCAAGAATTATGGAGTAATTGATGTTGTTTTG CATGTACTGGAAGAAAAAGGAAGTAGTGATTAGATTTAATTTTAATAGACATAT
>Gm16_3 1461554	TGCAGAAAATGAAGGAAGATGTCATGCTGTTGCAACCTGCTTAACCAGGAAATTATCTCTTC CACCCTCTACATTTCACTCTTTTCTTGCAACCATCATCCTCCAAAGTGACCCCAACCT ACTGGTTCAAGTAACAACCTTGATACCTGTTACACCGCGGCTTGACGAAGCCGTG[A/T]CGT GACAAGGGACAGAGTAAGAACTGGAATTTTGTGAAATTGCAATGGAGTCCTAGTCTAAA CTATAAGATAGAATAAGACTAGGGAATTTTCTGTGGGATAGTTTATAGCATGTTATTGCTAT TTTGCCCTTTGAATTTTGTATGGGTGCAATTTATCTCTTCTCTAGTTTTT
>Gm16_3 1475163	TATAAGTGTTGTTTTCAAATATTAAGGAGTTGTGGGGTGCGAGCGTGATTTTCTCGTTATAA AAATAAGAAGGAAGAGTGATTCAAATCTGAAGCTGAACCCTAACTAGCTGAGCAGACGGAG GAGAACTCGACTGAAGCTGCCTCTGCGACGCCGTTTCGATTCAGCCTCAGACATTT[G/A]TG TAAGTTCATCGTTTCACACTTTACACAAATACCCTATTCTTCGCATACTTCGTCGCTGCAACC CCGAACAACGATAACCTAATCAATTTGGCTTCGCCCCCTCTTTTTCTCTTCACAAACAATAA CAATTCACGTGTGATTGATTTCTCCAGTGTTTTACGAGTTTCAAAGATGCTCG

>Gm16_3 1476359	CATTTATCAAAAGCCCAGGTTAGCAATGCAAAACAGTGAAAGTGCTGCTTGCTGCTGTAG GTTGCAGTCATGGTGCAAGAAAATGGCAAGAGGAAGAAACCAACATTCTGGTGACTGGCA CACCGGGGACAGGAAAGACAACCTGTGTGCACTGCTCTAGCTGAAGCCACCCAGCTCT[C/T] TCACATCAATGTCGGAGAATTAGTCAAAGAAAAGAACTTGCATGATGGATGGGATGATGAG CTTGATTGTTACCTTCTTAATGAAGACTTGGTAAGCTATCAATTTATATAATTTGGATCATT CTATATGCTTGTGTTTGAGATTTAGCTCCTCTAAAGTAAAATGAGGAGATGTGTAA
>Gm16_3 1479958	TAGGACTTGTAGTTGGGGACACCATGAGACTATTAACCCCTTCTCTGATGTGTCAGCAAAC CCTTTGGAAGCTTTCCCTTATCACAATCTCTAATCACCCACATGAAATAACTTCCACTATCTC CCAAACCCCAAGCCAGTTCCTCAGTTTGCTCCTCATTAAGCCCTGCCATGCTCC[T/C]GAAA GAGACATACACAACCTGACCCTTTTGTTTTTCATCAAGCCATTTGATGCAAGCTTCTGAGTT TGGATTATACATGTTAACACCATAGTCCTTGTCTGCTTGGAGTCTTTTGCCAAGTAGATAG ATGGCAGGCATGGTCCTATTGGCTTTAATGGCCAAATCTTCACCAGCCAATC
>Gm16_3 1481163	ATAGCCATCTGAGATGCTCTCAACTTCAATGGAAGTGAAGTTTTTGTCTCATGTTCTTCCA ATTGGACACAACAGTTACAAGTGTACCTTCACTCCTCTCTGCACCAACGCTTGGAGA GAAGCATAGGATTGATGTGTCCTTGGGCTGGATAAGGTAGGACAAGACAGTGAG[G/T]TGC ATGGTTTTTCTTTGCTCTTTCTGACTCCAACGTTTGTACTCCTTCCCCACACTGC ATATGTGTTTCTGCTTTCTGTTACATGTGTATAAATAGAAGAGAAATGCAATTTTACACCA AAATCAACAATAAGAGAAAGAAAGGAGAAAAAATGTGATGTAATAAGTG
>Gm16_3 1506333	TAAGTAACGATAACAGGGGCCACAAGAGAGACGTTAACTAGCAGCTTCATTTGTCGCTACA AAATATTTGAGGTCACCAATTGAGGACACTAGAACCATAACAAATTTCAAATGCGCATACAA AATTTTCTTCCACTCTTCAAACACGCGAATTCTAATCTGAACCTTCAACACTCCG[T/G]GGA ACAAAAATGCTACTCCTTCCCGTGCTTTCTCTCTCCTCCTTCCCCTCTCTCGCTTCGAA CCGTTATCTCCAATTTGACCTCGTCGGAGCCGCGCGTCGTCGGATGCGCCGCCCGGA GTCCGGTGGCGATGAGTTCACGGCGAAGTCGGGGTACCTGTTGAGCTGAGCGCC
>Gm16_3 1516052	TCGTAATTTGCAATATGGGGATCTTTTTGGATTTTACTTGTTTTATTATACCTGACTTTT CAATCGTTTCAAGCAATAGGTATATTAACCTTGAACGCTACATGGACAATCATTTTACTGTTT GTTTATCCTAATAAAAAAATTGTACTTTTCAATGACATGTAGGAAAGC[C/A]GACAGGC GAGTCTGACAAGAATCAAATACTGTACATGGAGGAGTTGTCGTTAGTTTTAAATCAAGTTG AATCCATTCAAGACATTCTTCTATAATTTCTGTCTTCTGAGGTAAATCTAAGCATTAGCA TTTTATTTAAGAAATGTGGCTGTTTACCCATTTTTTCAATTAACCTGTA
>Gm16_3 1516059	TTGCAATATGGGGATCTTTTTGGATTTTACTTGTTTTATTATACCTGACTTTT TCAGCAATAGGTATATTAACCTTGAACGCTACATGGACAATCATTTTACTGTTTTGTTTATC CTAATAAAAAAATTGTACTTTTCAATGACATGTAGGAAAGCAGACAGG[T/A]GAGTCCT GACAAGAATCAAATACTGTACATGGAGGAGTTGTCGTTAGTTTTAAATCAAGTTGAATCCAT TCAAGACATTCTTCTATAATTTCTGTCTTCTGAGGTAAATCTAAGCATTAGCATTTTATTT AAGAAATGTGGCTGTTTACCCATTTTTTCAATTAACCTGTATAAGTCA
>Gm16_3 1520234	TTGATCATCAATGGGTTTTCAATTGAACGGAAGATCTCACTTCTCAGTTGCTGTTTATGCAT GTATATTTTTGGATAAAATTCAAAATTTGAAGATGTGGATCTAGTGTGAGAAAATTAATGAAT TGTGTGCTGTTTTCAACCGGTGTCAATAACATGGACCAATGGGAATGTTAGCC[C/G]TAGA TTGAAGTCCATGTTTTGTAACAAAAGCTTGCAGGGAAGTATCATTTAGCCTAACCTTGTA AGGGGCAAACTAAATTATGGATCTGATGAATTGGTGAGCAAGTGTAAGGAAAAAATATATG CAGTGGGAACAAAAATTGCGTACGGGACAAAAGGCACCAAGCATCATTGTTA
>Gm16_3 1520242	CAATGGGTTTTCAATTGAACGGAAGATCTCACTTCTCAGTTGCTGTTTATGCATGTATATTT TGGATAAAATTCAAAATTTGAAGATGTGGATCTAGTGTGAGAAAATTAATGAATTGTGTGCT GTTTTCAACCGGTGTCAATAACATGGACCAATGGGAATGTTAGCCATAGATTG[G/A]AGTC CATGTTTTGTAACAAAAGCTTGCAGGGAAGTATCATTTAGCCTAACCTTGTAAGGGGCAA ACTAAATTATGGATCTGATGAATTTGGTGAGCAAGTGTAAGGAAAAAATATATGCAGTGGGAA CAAAAATTGCGTACGGGACAAAAGGCACCAAGCATCATTGTTTACTTCTCTA
>Gm16_3 1525931	ATTATTGATCAGATAGAGGAACCTATTCAATGACTGTGTTCTTTGGAATAATGAATGAATCA TGTGGTGTGTAATCATGAGTACCCCAACCAATCAACAAATTGGAATCCATGTAATCA AACAAGGAAGCAACTTGGAGGATATCCAATTTACCAATCCACTATTGTCCAAAGA[G/T]GAC CCAGATTTTGCATACAACATGATGATTCAAGGAAACGAAAAAGACAAGAGTGACTGTCTTC TCCCGGAGATGAAGTATCTTCCCGTTGATTGATAATAAATTTTGTGGTTTTAGGACACCTT GTTTTGATTATATATTGTGCAATGTTATGTATTCCCAATATAATTACAGGAGG
>Gm16_3 1675129	ATTAACCGTGGCACAGCAAGATTTGAGTCCAGATTGTTAACCATTGTAGCCACTTTGTTAG CATCACCCATCCAATTGACAATAATTTGTTATCAACTAGTACATTACATCTTTTCAAGTGT TGATAAGAAAGTTTAGAATCACTAGGTATTGAGTAATGCTGTTTGTATCATATG[C/T]AAGGT

	GACATTCCTCAAATGTCAATATATTCCTGAAGAACACTTCTGAATTGTCAGTTATATTCAAGA TTGGCATTGTCAACACACCCCTCATCGGAATAGGTCATGTCAAGTAAGCATTCAATTTTCATTT GGACTGACCTTGAATTTAAGACCTGCCTCCCGTAGTTGGCTTGCACTGTA
>Gm16_3 1708589	AATTGACAAATATCTTTTATATTTAATTTTTACCTTAATAAGAAGAAAAGGACATAGGAAGG AATCAACACAATGCATGTTATGCCACAATTGCTAGAAAACGATACAAAGAAATCTGGCAGTA GAAAAAACAAAACTCCCTGTCAACCCCTTAAATCCCTTACATACCAATTACAT[G/A]TATCA CTCCAAAATTCCTACACCATGCCTTCACACAAAACTCCCTCGTGCCTATATATGAAACAAAG CCATTGATGCAAAAAATATGAAAAGAAGAAAAGGCAAAGGGAAAAATATCTTACATACAATGC ACACCTCAATCACATAATGTACATATACAGGATATAGCCAATCAAACAAAC
>Gm16_3 1708884	CTTACATACAATGCACACCTCAATCACATAATGTACATATACAGGATATAGCCAATCAAACA AACTCTAAATCTTATTGCTAAAAGGAAAAATCATATTACAAAAATTCAGAAAAAAAATGA ATGAATCTAATTCTAGTAGTGGCAATGTGAGTGTTAGAAGTTGCAGCAGCTATC[G/A]TTGAT GCTCCTGAAATTCTGCAAGTTAATCTCCTTTACAACCTCATGATGTATCCCTGCACCAGAAAAG TAGAATTAGATAATATATTTTATTACATCAGGAGCATGATTTTCAGAGATGTACAAATATTCAAA TTGATAATATTGAGACCTAGACAAGTAATAGTTGCTGCAATCCGGGAGAT
>Gm16_3 1709663	ATGCATGCTGACAAGTTTATACACGATATAAGAAGCAATTCACCTGTCCAGTTAAGCAGCTA AATTCCAATTACTTAAATTTGTCATGACTTGTTCAATCTCTGCATCAGTGAAAGTTA GGGCTCGCTGTAGCTTCCTAGGAGAAAATGTCTCAACTTCAACCATATATTTGG[G/C]AGTC ACAGGTCTGTGATCAGACAGTTTGAGCTCTGCCCTTCTGTAACCTCAATAATCTCATTCCCTT GCCATATGAAAGAATACGATCACACCTTCAAGGTGATTACAAACAAGCAGAAAAATTCATCAA GGGTAAGACATGTATCAAATACCATACCACTAGTTCATAAACCATGAAATGG
>Gm16_3 1790078	CTTTCATACCTTACCCTGCATCCCTCTCCCTCTTCTCCTTTTCTCCATAATTATAAAAAATCG CACCCAAACACCCCTTCCCTCTGCCACCCAACCAATCCTCGATCTCGCCGAACAACCCC ACCCCTTTCAAATCCGATTTTCCGATGCCGAAGAAGCGAAAAGTCTATCGCCACCA[G/A]CCT CGACGAGGTGGATCGAACCCTCTACGCGTCGTTTTGCACCGCCGCAACTCCCTCTCGCA GCTCTACACGCACTCCATGAACCACCAGAAAGCTCTCCTTCAACGCCGGGGAACGACACGC CCTCGTAATATAATAAACCTGCCCTCCTTTTTATTTTCTTTCTTTGTTTATTCTC
>Gm16_3 1791873	AATGTTGTAAATGTAGTGTCAAATGTTCCATTTGTGATGCTTAATCTTTAAAATTTATTGATAT TTCTTTGTTTTTGGTTGTAAGTATGTGAAAGTTCATTCCCTCTTGCAGAATGAGCTGGATTATT GTGGAGAAGAACCATCTATGTCACCTAGAGCACCCTGCAACACCAGTCACA[T/A]GCAGC ATTGCATGTCCCTAGTTTTCCAGTCACTTCAGCATCTTCTGGTCAAACAATAGCTGCACAAG GACTCCGCTCGGACCACTGTGAAAATCAATCAAAGAATTCTGTGTTTTCAAATGCTTTGTCA AGTCCAGTACGTGGAAGCCTTCAACATTATCAAATTGGTGAGGGAGGATGC
>Gm16_3 1792317	GTGAGCTCCAATGATTCTGCCATGGATATGCATGCAGACTAGTTCTGTCTATTGGTTTATTA ATTGGGAATTCTTAACTCACCAGCCTTTGCAACAGCTACAAAAGAAGCTAACAACTTCGGA ATCCGTATCTGGGTTTGTCTATTGGATCCTGCAAGAAACGACAGGAGGATGGAA[G/A]TAA TTCTGTTAACCTTTTACTATGCTTACGTCAACCGAGTGATTGGTTTGGGTCTTAAAGTGGTT TTTGTAAGTTTCTATTCAATAGCTTTAAAGCATTTTATTTTGTACCTTAAATTGTTTTACCC CTTCGCCCTTCACCCAAAAGACTAATTTTTAAATGTAGGGGTTAATTCT
>Gm16_3 1799191	TCGTTTACTGGACTAGTTGACACTTGACAGCCATTAGTTTTATATATGATGTTGCAAAACATT TTATTATTCGTGGATGAGCTGGATGTAATGGAAATTTCTAACCTTGTTACCTAGCCATATATT TATGTTGTTCCCTTTTCTATTAAATTGCAGCGAAGTGGATGATGAGTCAGTTG[A/G]TGAATT TGACAATGGGAATCTCAAACTTGTTCCCTGTGCTATTATGTAAATTGGCATTCCAAAGGATG ACACAGATCCATTTATGAGGCGCCAAGTTGTTGATTGCTAACATATTCATCACTCAGCTTG AGGAAAAGGAGCAGATATTTTTCAGTCTAGTGGAGATTTATACACCATT
>Gm16_3 1827884	GAAACAACAATCCCCAAAATAGAACACACCCATATCCTTAGCAATTAACAACACCACACAA GAAGAATACTACATAACTTACAACTTTTGTCCATTCAAACTGATCAACCAATTGCACCA CCTCTTCCACCACAGACGTCAACAAAGACTTCCCTAGCATTACACATTCAGCTT[G/A]CCA CGAATTTACCTTGATTTCCAGTGAGCACATCCAATTGACTCAACTTTATAGTAGCATCCAC AACTTCTCAAAGAACAACGTTTGGTTCAACGCAAATGCATTGACCAACCCCTTAGTCCTCT TATCATTGAGCAAGTCTGGTCAGAGGTGAACACCCCTTGCGGTTTCATTAGG
>Gm16_3 1827991	ATCAACCAATTGCACCACCTCTTCCACCACAGACGTCAACAAAGACTTCTAGCATTACCA CATTGCATTGCCACGAATTTACCTTGATTTCCAGTGAGCACATCCAATTGACTCAACTTT ATAGTAGCATCCACAACTTCTCAAAGAACAACGTTTGGTTCAACGCAAATGCAT[C/G]GAC CAACCCCTTAGTCCTCTTATCATTGAGCAAGTCTGGTCAGAGGTGAACACCCCTTGCGCG TTCATTAGGTCAAGGTAGTATTTGTTGTGCAACACAGTCGGGGTTCTGATGTCCAAGTTGAC GGTGTGCGCGAGTTTGCCTCGGGCAAGTGGATTGTAGTTGCTTGGCTAGGGT

>Gm16_3 1828137	CAAAGAACAACGTTTGGTTCAACGCAAATGCATTGACCAACCCCTTAGTCCTCTTATCATTG AGCAAGTCCTGGTCAGAGGTGAACACCCCTTGGCGTTTCATTAGGTCAAGGTAGTATTTGT TGTCGAACACAGTCGGGGTTCTGATGTCCAAGTTGACGGTGTTGCCGGAGTTTGCG[C/G]C GGGGCAAGTGGATTGTAGTTGCTTGGCTAGGGTTTTGTCCATGTTGGGGTCGAGAGGGGA GAGCCTGTTGAAGAATGTGCCGCAATGGGCACGACCAAAGGTGTGTGCGCCGGATAAGGC GACAACGTCGGTGACATCAAAGTTTTTGGCGGCGAAAAGCGTCAAGGGTCACGCCGGTGG
>Gm16_3 1829738	ATATCACAAAGAAAAGCTTCAAACTATAGTTAGTAAGAGCAGAGGGTTTTGAGAACCAAAA TAGTAGTGCAAGTAAAGGTAAGAGGGATGAAGAGATTACAAGGAAAACCTGAATCACGTGCT GCCAAAACAGTGATATCTGCGCAAGAAACGATCCTTCCACACTCCTTATGGATAAT[C/G]GC TCTGATGTCATCAATGGTTTGCAAGGCCTCAGTTCTGATGCCTCCGTTGGCTGGTTGGTCT CTTTCACCTTGGACTCCCATCTAGCAGCAATGATCCATCACATCCCTATACAAATGCAAATTA AGTTAGAGGTAATATCTTTTTTCTATATATATATATGCATGCATATATATTAACC
>Gm16_3 1840753	GTTGGGGGACTCGGCATTGGGACACGTAGCTATCAAATTGTTGTTGAAGTTTGGGTCGATT GGTGGGTCCGTTTCGATGGTTCTGTTCACTAGGGAGGGACAGTGGGCACGACCGTAGGTG TGTGCGCCGGAGAGAGCGACCACATCGGTGGCATCGAAACCTCTATTTCCGAATCCTC[A/G ]CAGAAGATCGTCGGTTCGGAAGAATGGTGCCGGTAGGTTGTCCGGCGCGGTGGCGTTTG GTCCTAGGCCGTCTTTTCTTCCAGTGGCACGTCAAATCAGGGCCTCCTAACTGTTCAAA TTACAAAGAAGTTACCAAAAGGAACAAACGAGAGATCAAGCATTATGTTTAAATGTGTTT
>Gm16_3 1840819	GTCCGTTTCGATGGTCTGTTCACTAGGGAGGGACAGTGGGCACGACCGTAGGTGTGTGC GCCGGAGAGAGCGACCACATCGGTGGCATCGAAACCTCTATTTCCGAATCCTCTCAGAAAG ATCGTCGGTTCGGAAGAATGGTGCCGGTAGGTTGTCCGGCGCGGTGGCGTTTGGTCCTA[ G/A]GCCGTCTTTTCTTCCAGTGGCACGTCAAATCAGGGCCTCCTAACTGTTCAAATTACA AAGAAGTTACCAAAAGGAACAAACGAGAGATCAAGCATTATGTTTAAATGTGTTTTTAATCT CATTATAATGTTATAATTTGAGTGAAAATTATCTCTAATTTATATATTATAAATGATTT
>Gm16_3 1842815	GAGCTACTCCACTGTCCTTCTCGAAGACATCCTCAAGATGCTTCCTTATAATCCTCTCAAGC TTGGGACATGTCAACAAATAGTAATTCATGATAGTCCAGGGACTAGTTTCCAGAGGAAAA CATGGATTTGAGAAGCCAATACAAGGGAAGAAATCAAGAGCAAAGAGTGAAACAAA[G/T]GA CTAACCTAGCCATGTTTGATTAATTTGCTATGAGTGTGACTTGAGATAACCAAACTTAGA AGAGTTATTTATAGATTCAATATTGAGGTAACGTTCTTCGGTTGCTTCCCACTTCCACCCA CTAGTCTTCTACGTATAATATTTAAACGTGGCGCAAGCGTTTTTTGAGATTTG
>Gm16_3 1885123	TGTCTTGGCCGATTTTCTCTCTACTCTGTACACAGAGACAACCATGCCGTGGCTCCAGGT CTTCTTATTTTCTTAACATTATGTTATTCCAGATGCCTCAAAAAATGGTGTTGCACAAGAGAC AATTCAAATACTATCCGCAGAGCTAGCAAGGTGATGACGGAGCAAGAGGCTCGA[A/C]GG ATTCTTGGTGTTACAGAGGAAACTCCCTGGGAGGAGATTATCAAGGTTAGTATATCATCTTT GTTGTGGGTATATTACCCATTTGTTTTCAATTTAGCACTTCTAAAGTGGGAATGCATTTTGG AGGATAAAGCACACTGTTATAGCCTTCAATTGGAATCAATAGCTAACATAT
>Gm17_4 966500	CAGCAACACCAAGAAACATAAAACAGAAGTAAGTGTCAAATCCACCTCCAACGCCTACAGC TACACTGGGACCTGGTGCAAAGAATGAAAATGGAGACCATCCCCAGCCACCATAACCATAG CCATAGCCACCAACAAGCGGAGGTGCCACTCGAGGATTGATATAGATGTTGGTCTCTG[T/C]T AATTGTCATAAAATTCAGAAACAAGATTCCCAAGTTAATTAATCAGTTGATATATATGTTAT ATGTTTCATATGATCCATCCAATTATAAAAGATATTTCAATTCCTTTGATATGCATGATCTCTG GGGATACTCCAGGGTACGAATATCACATATCAGCCATAGTTAAATAGAATAAA
>Gm17_1 0335675	GCGGCGACGGGGCGTGGTGTGGAGGCATTTCTCGGCGCGAGAACGTGAGGGTGGT GAAATTCGGGACTCGACGTGGAAGTGAGATTGTTGGGGTGTATATTGCGCACCCCATGGC CAAATCCACCATCCTTTACTCTCATGGGAATGCGGCTGATATCGGACACATGCTTGAGCT[T/ C]TATGTGGATCTAAGTACCCACTTGCCTGTTAATCTTTTTGGGTATTTCAAGTTTTCTCCTTT CTTTCATTTTTCTATTTCTCATCTTATGTACATACATACATGATACATGGTACATGCATCACA TTGATCTGATTTTGCTGAATTGCGTTTGATGAAAAGTCTAGTGAAGCAGGGATG
>Gm17_1 1391400	TAGCATGCAGCTCTGTTACTATAAGCCTAAAAAGAAAAGGGATATCTAAACAGATCAGGATTT GCATTCATATTGAACACGAGGATTACTGCTATTCAACAGGTATTATTACCTTAGCATCT TTTGGGTTCTTTTTATAGCTTCTGTATAATGCTTCGTAGCCTCAGGATACTTC[G/C]GCTGC TTGAATAATTCAATTCCTGCATTTTCAAAGTTTCCAGTATTTAGAAAAGTTAGTTAATATTTAA GCAGAGATAAGAACTTATAGTCATCAAGATACATAGTTAAAGAGTGTTCCTTTTACAGGAG AAAAGATGCTTGACGAGATGGAGAACCCTAAGATTGTAGACAAAAAGAT
>Gm18_2 1030712	TGGATATAGCTCTCAGGGTTCTGGATCCACACTTATTATGCCATTTCTTGATAACCAGCTGA AGTCTCCTAGCCCACTCCTGTTACCTGCGCAGGTGAGGTTCTGAAATGGGTGTTTTGCTTG AAATATTTGATTCTTCAAGTTTATTGACTGACACCTACATTAATTAATGATTTCT[T/C]TCAG

	GATGCTGTTACTCCATTGTCTGAAGCAGAAGCAGTTGATCTGGTCAAACTGTTTTGCATC TGCAACTGAGAGAGATATACACTGTAAGTTTGTGCTAAATACATAATTACCAATTGACTG TAGGATGAACTTGTAACCTTACTGTGGATGCTTTTACAGGGAGATAAGGTTGA
>Gm18_4 1433527	AGGGTGAAATGAGCTTTATTTTTCTCGAGTTTTCTAACAGAAAAGTCAACCCGGTCCCTTA ATTGACGAGAATTCGTGAACAAAAGCAGTAAAATTTCAACGGATCAAAGGATATGAGTCATT ATCGCAGATCGATTCTTCGAGAACGAAACGAGCTCGGAGACGAGGGGAAACCCTG[T/C]AA AATTGAAATCGGGGAAAATAACAATTACAATTAACGAATTGAAGCAAAAATCAGAGAGAAGAG CAAAACAGAAAAGAGAACGAATGGAAGAGTACCTGATCGAGAGAAAGCAGAGAGCAAGATT GGTGCGCAAAGTATGAAAACGTGGAAGGGGCGAAACGACAAAATAAGGGCGGTTT
>Gm18_5 4168590	CCGCTCGTAGCTGTCTTCCTCAACTTCTGCATTTCTATGGCCTCAAACCTCTTCCTCTCCC TAATCCTAGCATTCTCCGACACGCCAAGCTGACAGCCAAGGCATCAAAATCTCATACCA AAAGAGAGTAACCAACAAGACAAAGACACAAGCAAGACGAGGCGAAGTAATCTCGA[G/C]T CGAAAAGCGGGCGGCAGAAAGCTGAGAGAGCGCCAGAGACCAATTAACGGAATAACCAAC CAAGGCAGCATTGTAGCAACCCGCGGAGACCATTCTGAATCACACACAGTATACACATTA TCCTCTAACCTAACCTCTTTCCAGTCCCAATATATCCTATCTTTTTTCCCTCCCA
>Gm18_5 9757318	AAACTCACAGATTTTGGCATTAGCGTACTGGGACCGCGTTTTATGTCAAAACCAAAACCAAT TAAAGTAGATTCTGTTGTGGGTAATGTTTTATTTTATGAGGTTACTACTACAATGATTTTTTG TCAGATAAATTAAGCTAATACTTTTACTTATAGGTACTTTTGGCTACCTGGCTA[T/G]GAGC ATCTCACGGACCTCACCGTCACAGATAAACTGATGTTTACTCATTTGGTATGGTTCTACTT GAAGTTGTGTGTGGAAGGAAATATGTAACAACAGAAGTTGAGAAGCCTGTTGAGGAGAAAA TTGATCCTAATATCAAAGGAAAGATTGCACCAGAGTGTTGGCAAGTCTTCAT
>Gm18_5 9757325	CAGATTTTGGCATTAGCGTACTGGGACCGCGTTTTATGTCAAAACCAAAACCAATTAAGTA GATTCTGTTGTGGGTAATGTTTTATTTTATGAGGTTACTACTACAATGATTTTTTGTGAGATA AATTAAGCTAATACTTTTACTTATAGGTACTTTTGGCTACCTGGCTATGGAGCA[A/C]CTCAC GGACCTCACCGTCACAGATAAACTGATGTTTACTCATTTGGTATGGTTCTACTTGAAGTTG TGTGTGGAAGGAAATATGTAACAACAGAAGTTGAGAAGCCTGTTGAGGAGAAAATTGATCC TAATATCAAAGGAAAGATTGCACCAGAGTGTTGGCAAGTCTTCATAGATATC
>Gm19_6 362210	CCTTCTTCTCTTCCCTCCCAACCATCAGATACCCAACTTGGCTGCTTCATATTCTCCACCTCA GCTATATATATATATATATATATATATATATATATATATATATATATATATATATATATA TATATAAATCAATTGAACCAAAACCAATCTCACAATAAAAATCACAATCCT[T/C]ACCTTCAAC TTTTCCCTCAGTAAATCAGCCTCCTGCAATTTCTCCTTTATTTTATTCTCACAATCTGATATTG CATTTTTCGAACGCAGCATTCTCTTCTTCAACGTCACCTCTCTAAGTTGGGCCTGCACCACA ATTGCATTTAAGTGTCCCAAGTTCCAATTGTATGAATTGAATCCC
>Gm19_4 1528666	ATCGTCATGTACCCCGTTTGGAAAAGATGTCTCACCTCACCATACTTGCAAGTCAAAATCAG AATCAAGGGAACAAAACGGAATCGCGAGAACTCTATCATGTGATTTTTATTAACCGCTATT CACTCCGACAACAGATAAAAATAAAGCTTCAGCCTTCTGCACCTTTTTTTTTTTT[T/C]CCTTC TTTAATCAAAGCCTAAACTAACCTTGGATACAAGTCACAGCGTCAAAATATTAAATAAAGAA ACGCCGTTAGCCAGAGGCTTACGGCGGAAGCTTGCACGGACTAGGTGGAAGACTTGTCTT CCCAGTAGCAACAAGACGTTGCACGAAGCTCTCGATTCTGTTACCGTTAGTC
>Gm19_4 9218118	AACCCCTCCAACCACCATAGCTGCCGTCCTTGGGCACAATGCAAAAGTTGTGGAGCACATC AGAGTGGATTGCTTCCATTGTCTAGTTTTTATGAATAAGTCTCAATTGAGACAGTGTTGCG TTCCCTAAGTTTGATTACATCTTGCACGGATCACCTTGTAGCATATGTCACTAT[G/C]AAG ATATTCCTCAACAATCAATGCCATGCACAGAACTTTATAGAAGTCTGAGGTTCTGGGAGTT CATACTTTTGTGTTGGTCACAGTATTGTACACATAATATGTCTTTGGATGGAAACCAGAAAGA ATAATTCCTTTGGAAGTATCAATGAAGTACCCAAGATGTTTGAGGATTCCAGA
>Gm20_3 7990145	TGATGTAGGTGTTCCCTACCCACCGGATTCCACCCCGCTCCAAATCCGACGTCACGCG CTGGCAGAGCTTCGTCCGCGAGCGTCAGCGCCACGCGCTCTTCTGCTTCGCGGCGCGCG CTCGCCGCGCCTTCGCGGACGACTTCGCGCGGATTCTGCTGAGTCAGTGCCGCGACTCGG [T/C]TGAGTCGTGCCGCGCCGTGAACTGCACCGGGACGCGGTGCTCCAACGGCACGTCGG CGATTCTGGAAGCTTTTCTGGACTCGGATTTCTGCTTGCAGCCGAGGGGGGACAGCTTCAC GCGCCGGTCCATTTTCGACTGCATGGTGGCCGGTTCGATTCCGGTTTTTTTTCTGGCGGCGA AC
>Gm20_4 4848539	AATGCCTCACCACAGACACAGTTACAACACTGTTTCAATAAGTGACTGGTCTCCGAAACAAA GGACCCATCTTCGAAATTTGAACTCTGCCTTCTCAATTAAGAATTTAAACTTGTCAAATTAAG GGGGGGAAGTCAAGAAGTTCGATGGTGGTTGCGGAAGCGAATATAATATAATCAA[T/C]TGG GGGTATAATTTGTTCTGATAGGCTGAGTCCCCACAACGGAAGCAACGAACCTTCTCTCGTC AAGAAACAAATCCTCGGTAAACCTAAACGCAGCGTGCAAGCCAACGAGAACCACCGAGAC

	GATTAACGAGAGCAGGACGTTGAGGCCGACGTGAGTGGAGACGAGCGAAACGACAG
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**Supplemental Table S5.** SNPs identified by Method 2 on the seed protein NIL pair. This table contains 387 SNPs identified using Method 2 on the fourteen Illumina libraries of Williams82/G. soja seed-protein NIL experimental data. The file contains a unique SNP identifier that includes the genomic location. The sequence surrounding each SNP based on the soybean Glyma1.01 genome assembly is also provided.

SNP	[W82/Gsoja]
>Gm01_244437	TGTCGTGTTGCTTCTCGTTGTTATTTTTTTTTTTGGGCCAAAGCTGCTAACCTTCTCAATAGACCTCTTCCTCCTCTCCCCGGTCTGGAGATCTGGAGGAACTATTGCCAGGGTTGTTGATGC ACTGCAAATTTGGTTGAATCTAGCCTTGCTGTTGCACATGACATAGCAATTGAG[C/T]AGAA TTTTCTGCTTTGCCTCCAGGTTGTTGGTGTATTGTGGGTAATATCTTACATCGATAGTTTGTT CAACTTTTTGACTTTGATCTATATCGGTGTTCTTCTCGCTTGCTCTCCTTGTTGTATGA CAAGAATCAGAACAAATTTCAATTTTCATGGCTAGAGTAAGTTTCTCCATG
>Gm01_526411	TACATACACTACTGCTAAATAGAAAGAAATGCAGACCTGATTGATCCCCGATCTTGAATAC TGTCAACTATCTTTTTGGTGTCATGTCCACCTCGTGATGTTCCGAGGACAGTTCCCCACG CTTATGTATATCATTACACACTTTTAGGGGTTAAAGTGATTGTATTGCGAGCATAGA[G/T]ACC CCTATATCCTCCCTTCACATCAAAACATTTTCATCAGTTGATGAGACCAATACAAAATAAG TGTGCTAGCAAAATGTTAGTTATTGTAGATTTACACTTCTACACACAAGGGAGAACAAAAGA TGGTGAAAATTAAGTCCAGGATAAATAACATATATAAGTGACACCTCATAAGT
>Gm01_2450043	ACTCTGGAACAGGTTTCGCAAGTTGCTTAAGGATTTCTGATGATGACGAAGATGTATGCTAC TTTCATTTTCAATAAGATGCGTTTTGTGTTTGTCTTGCTCTAATACTAGTGATGTAGTTCT GGAATGGTAATTTTGTATTCTATTAATCAATGAGAAGCACACATTATTCTCCGA[T/C]TAAA CTTAGATAATCTTATTTTTTCAATTTGATCAAAAGGCTAATTTTATTTATCATATATGACTCTCA Ttagtctaataaatttttttaagcacataaaaaattaaatataaattttttattaaataaatcg TGTTCAATTTATGAAAATGTATTAGAGTCTTACTGCACCAAATTTT
>Gm01_2524255	TTGCAAGCGATAGAAATTGATAAGGCCATAATTGATCTCTTAGTAGTAGAGAAGGTGAGGT TGCTGTTCCACATAGAGCATATTGTCTTGTGTTGAATTGATAATTGGTGTTAAGGTCATCT TAGTGAAGTGTCTGACTTTTTGTGCTATGCAATTTTCATTTTGCAGTGAGAGTTC[G/A]GAA GTGTGTTGAATTGATAAGAGATTTGTCCTTGATCAATGTTGCTGCTAGTTTAGGCAGCAAGG TACCATATGTGCCCCAAAGTCCAATTGGGGTGCTGGATGCTGGGTGCTTAAGTTATAAGAG TGATGAATTAAGTGTGATCATGTCCAAATTTTACATAATATTTCAAATCC
>Gm01_7695499	CAAAACAAGATATTGAAAAATAGAAGAATGCAGTATAAACTGTCAAATGGGGAACCCCTTTT TTCAAAGTATTCCATTGGTCATTTGTTGCTATGTTGAATGCTGGTGAGAACATAGAGAAGCA AAGTCCAGCAAAGAATGTTAGAGCCAATCCAATCAAAGTGCTCTTCCCAAAAAC[C/T]GCA GCAAAATTCAAGTGTGAGATATAATCATTTTCAAGTAAAGGCCAAGGATTTGCCTAACAAGGT TGGTTAATTGAATGCATGTTTAAAGTCATCATCCAATTACACATTAAATATATCATCATAAAGA GAAATTTCTAGCCACACTCTCTTTGATACTTTCTTTCTTATACTCTTTGA
>Gm01_8945342	TTTTATTCAAGTTATAATAAGTAGATTGAAGATTGTAATTGAAGGTTATAATAAATTTTTT TAATATATATATATATATATATATATATGCTTTTCTTGCTCCCAATTAGTGTTTCCTGGTTCA TATTCTTGACTTTTCTTAATGATTACGATGTTGGTGGTTGAGTTGTCATTG[A/G]AGAGTTTAT AAACCTTGCTCTCGTCAGAATCCAGTGAAATTTGTAATAGAGAGGATGTTGAGAGATAGTATA CGTTGAAATTTTTGGGTTTTAGGCCTTGAATTAATGATAGGTTTCTTATTCTTAGACATTTCA TATAATATTCTTAGACATTTCAACTAAATTTGTGTTGGTGTGTAGC
>Gm01_8945369	GAAGATTGTAATTGAAGGTTATAATAAATAATTTTTTAAATATATATATATATATATATAT GCTTTTCTTGCTCCCAATTAGTGTTTCTTGTTTCATATTCTTGACTTTTCTTAATGATTACG ATGTTGGTGGTTGAGTTGTCATTGTAGAGTTTATAAACCTTGTCTCGTCAG[G/C]ATCCAGT GAAATTTGTAATAGAGAGGATGTTGAGAGATAGTATACGTTGAAATTTTTGGGTTTTAGGCC TTGAATTAATGATAGGTTTCTTATTCTTAGACATTTCAATAATATTCTTAGACATTTCAACTA AATTTGTGTTGGTGTGTAGCATACCATGCTTTAAGGTTTTTAATTTG
>Gm01_13799455	TTCAAAACAATCTTGTTTCAAAACAATGGCAGATAGATCCGTGTTTCGAGATGACTTTGGTAG ATAGAAGATAGGATATTGAGGGAAATAGGGTTGTAAAAAATGACAATTTTGAACGAAGA TTCAAATCGTCTCGGCATAGAGAGAGAATTTATGGCATCGAATTGAACTAGATCT[G/T]TG GCAACAGTGCAAGAAAATGAAAACAAAACGTGACAAATCGTGCAAGAAAGAAATTTAGTTT AAATATTTGCAAGTGAGAAAGAAAAGTTGAGAGAGGATCCGATTTTCAAATGATGTAATC

	TAGGATTACAGAAATTTTCTTACCTGGTTGGTTAGATTCCCGTGAAGTGAAG
>Gm01_25562169	ATGGATGGTGGGAAGGCATTGTTGTTCAAAGGACTCGGAATCTAATTGTCATGTTTATTTCCAGGTATGAATGTCTGCTCTTTCTATGTTAATTAGCTTAGGTTGGTAAGTGTCTTTTGGTACTTAGATTATTAGTTCGTTTTTTTTATTTTTGATTAATTCCTATGGTAATGTCCTT[T/A]GCAGGGGAAAAAGTGGTGTGCGTATTTGGTCCTGGTAAGTTGAGAGAGTCTCAAGATTGGGTGGGGAATGAGTGGGTATTTGTGAGGGAAAGACCTGATCTTGTGGCTTCTGTTTTATCTAGCTTGAAACAAAACAAAACCTCATGCAATCCAATTCCAATAACATCCAATCAACTGGA
>Gm01_27083133	TTGCTGCTGTGGCTGCAGTTGCTGCAACTGCGGAAGCTGCTGCTGTTGCTGTAGTTGTGGGAGTTGTTGAAGTTGAGGCAGCTGCTGCTGTTGCTGGAGTTGCGGAAGCTGCTGCTGCTGAGCTGTTGCTGCTGTTGTAGCTGAGGAAGTTGCTGCTGCTGTTGCAGCTGAGGAAGCT[T/G]CTGCTGCTGCTGCAGCTGAGGAAGTTGTTGTTGCTGCATGTGAGGGTAGCTGCTGCTGTTGTTGCAACTCGGAAGATGTTGCTGTTGGTTTTGCATCTGCTGGTGTGCTGTTGCTGCTGCTGACTCGAACAATTGCGGCTTAAATACGACCATATTCCTGCTTGCATTCAAGGCTTCGAGTAA
>Gm01_48157361	TGGCGGTAAAAAATACAAAATCACGATGAAATCAGTGATAATGACTACTCCAAACGTGGTTTACGAGTGAGGGACCCACCTGTAGTTTGTATTACCGAAGCCGACAACACTCTTCAGATCACTCTCATTCTCTGACACCGGGAAGCTCAAACGACGCCGTGTGTGTTGCTGTTGC[C/G]GCTTCTTCTCTCTCTCTTCTTCTTCTTCTTCTTCTTCTCACACATTGTTCTTTCTTCCATCATCAAAGGTGGAGCAGCTGAAGACGCTGGTGGGGAGGGAGCTGACAATGGGCTCTCACCATGGCCACTCGAAGGAGTTTCTGGACCTCATCAAGTCCATCGGCGAGGCCCGATCC
>Gm01_48157362	GGCGGTAAAAAATACAAAATCACGATGAAATCAGTGATAATGACTACTCCAAACGTGGTTTACGAGTGAGGGACCCACCTGTAGTTTGTATTACCGAAGCCGACAACACTCTTCAGATCATCTCTATTCTCTGACACCGGGAAGCTCAAACGACGCCGTGTGTGTTGCTGTTGCA[A/G]GCTTCTTCTCTCTCTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCTTCAAAGGTGGAGCAGCTGAAGACGCTGGTGGGGAGGGAGCTGACAATGGGCTCTCACCATGGCCACTCGAAGGAGTTTCTGGACCTCATCAAGTCCATCGGCGAGGCCCGATCCA
>Gm01_51613297	TGAGAAGTGACGCGTGTCTATGACAGCTGTATCCAAATCCCAAGCCAACCGCTTCCCTGGAAGAAGCGGTACCGATTGCCATAACGCTCGCTGACAAGTCCAAACCTAAGTTGCCAAACCGGTGTCTCACACCCTCTCCCTCTCAGATATAAGCGAGCAAACCATTCGAATGCT[A/G]CACACCACTACGCCACAGGAGGAGCTTCTGTTTCCATTTCTCTCACAAAAATCACTTATTATGAACACACGAAAAGACATGGACCGGATCAAAGGCCCATGGAGCCCGGAGGAGGACGAGGCTCTGCAGAAGCTGGTGGAGAAACACGGGCCAGGAAGTGGTCCCTCATCAGCAAGT
>Gm02_858015	TCCTCACTGAGTTTTGGTGTAGCCTGAAACAATATACATCAATTGCTGTCATCAGTTTCATCAGCTAGGATTTTAGAATCGACAATAGACCATAAAAAAACTTAGTTTTCTCCTCTTTGACGGAGTGTGTGTTGAGGGGAGAGGGGATTAAGATAAGATAGCTTAAACTAG[C/T]ACCCAAGTAACCAAGACTCTGTTGTCCACGTGGTAGTGATGATCAACGGGTTTTCTTCCAGTCAGAAGTTCCAGAAGGACAACGCCAAAACCTGTATACATCACTCTTAGCATTCAATTGTCCAGTCAATTGCATATCTGCATCCACAAGATTAGACAGATAGTAAATCAGCAGTCTGAGGC
>Gm02_1882327	CCATATGCTGACATAGCCTAGAAAAGTAAACCAAACAGATTACATAGTTCTCCATTCTTAATTTAAAAATAAAATGAAAGGAAATACCTTCCCAAGTTCTTCGTCCACTGCTTTTGCAATTGTGTTCTTCTGCTCAAAAACAGCATCAAGTTCCATCTTTGGAACAGTAGCTCTGATAA[G/T]TAAATAACAATAGGAGTAAATTAGTTTCAACCTGAAGATAAGATGCACACATAAAATTAACCTAAATTTCTTCTTCTAAGTAGATTCACAACACAAGCTCTAATTGAAATATTAACCATCAAAGACATAGGATTGTATCTGTGCTTTGGTATTGGTGAGTTTGTAGTAAGCATCTGATG
>Gm02_4060077	CCTTAATAGAAAAAAAACCTTAGGTATTTTGATTTTTATTTTATTTCTGGTGGTGATAAAGAAAAAGTTGCTAAGCTGTTTATTTATTTGTGAATTTTGCATTTGCAAGACTTTCAGCTTGGGGAGACTGGTAGAAGTTTGCCCTCATTGTCATCACATTTTTACCTACCTTGCATTGAT[T/C]AGTGGCTGATCAAGCATGGTTCTTGCCCATATGCAGAAGGGATCTGTAATTTTGTATCTGCAAAGTATACAAACCGTAAAAATAGCTAGGTAACTCTTTCATTTTTATATATAAGGGAATCTACTTTAGGAATGTACACTATGGATAAATGTAGTCGGAAGTGAAGCAACCATC
>Gm02_12048612	TGAAGCCAGGAAGCAGTGATTGCAACTCATGGAGCACTTCAGGTAACCTTCTATTGTAAAAAGCTGATAGCTTTGTTTATTTTCTCATCACAAATTTCTCTTGGTCTTTCTCGGATAGCTTTGTGGAAGCAACCTGCTGGGGGAATGTTGTTACAGAAAAATTTCTTGCTCCTAGA[G/A]CATATATTCTCTGCCAGAAAACAACAAATCAAATAAATCCCGTTAGTGATGTTATTGCTGATGAACTTTCTTGTTGACAAGTGACAACATGAATGCTAGCGAGTGTAATCATTTGAAAGAGAATTCATGATAAGCAGCGCATACATAGTCCACTTCTGTTTAACTAGTGTGATAA
>Gm02_1485198	GTTGGAGAAAGTGGGACGCTTGTGAAAATATTGAATTGAGCCTTGGATTTCGCCGCCGAATGGCCACCAAGAGCTTTTCTTGGTGGGGGCCAACGCACGCACCCGTTCTCTGCTTATTTTGA

3	AAATACTTCCCTCGTTCTGTTGATTA AAAAGCCTTTCTAACTTTCTTTTGGGGTCT[C/T]TAA TTAATTTCCCAAATAAAAAAGAAATCAAATTGACAAAGCCAAATTAATTAGTAGCATTAAATG TTAATGATGCATGATAGCAAGAAATTAAGGGTCTATATACTTGGAGCGCAGCTTGACAGCAC ATTCCAGCTGAATGCTGAGAGGACCGATTCTGGGTCTCGTCTCCAGGCCCGA
>Gm02_ 3734972 6	TTGACCGGATTACCTCTCCTCCTCATGCTAACTATTCATACTCTGTTCAAGCTAGTGTCAGT GCTGGAATTGATATGGTAACTTTTATCATCAATTTTAAGGAGTAAAGAAACTTATATAAAAT GTTTTTTCAGCCTGAGAATTTTGTTATTTCAATTAACAGATTATGGTTCCC[A/G]ACAAC ACACCGAGTTTCATTGATGAACCTAACCCATCAAGTAAAGAACAACATTATCTCAATGAGCAG GATTGATGATGCCGTGGCAAGAATCTTAAGAGTAAAATTTGTCATGGGCTTATTTGAAAATC CATATGCTGATCCAAGCCTGGTGAACCAACTGGGTAGCAAGGTATATTTTC
>Gm02_ 4224890 3	CAATCCCAGATGTCAATTTCTTATCCCTGCAACATAATTACATAGCAACCATCAAGCAGAT TTCAGACACAACATAATAAGCTTTCTATGTGGTATGCATTAGAAATGTTATAAATTTTATGAC ACTAAAATAATTACTCACTCACAGTCACTGCCCTAACATTTTTTTCTTCTTCT[G/C]ACCAGT TACCCTTAAACTGGGAAACCGGTCTCCCGTTTACGTTGGTCCCACCACTGAAGACCGAA GCCGAACCCACCGGTGGCGAACTCCATCGGCGACGCCACCTCACTGCGGTGTATGAAAC CAACCCGGCGCTATCAAACAACCTCCGGTAATTCAAATTGGAATCCCCGATAC
>Gm02_ 4752229 3	TTCTCATAACAAAGTAAATGAGTGAGTTTGATATTATCTGTCCAAAATATGAATTTTCTACA GTAACAGCACTTTTCATCTTCACTTGTGCATTTTCAGATTGGATGTTAGCTTGCATTCTCTGA ATCACTCTTCTGAGGTGAGGTGGATGTCAGTTTTTCAGCAGATAGGTCAGAAAGT[G/T]CTAC CATCTGAAATAACACAGCAACATAAGTAATAAAATTTGTTACAAATTATAATATTTAATTTTCT CATCCAAAATACTTGCTATAATGTGCATTGAAATTACATTTGATTTTACGCTGGGGCTCAT TGTTCTACTGCAAGCCATATTCCTGTGCATGGTATCTATTGCAACAAAGT
>Gm02_ 4775192 2	AAAATGGAATCAAATTCGGAAGAGGAACTGAATAGCAGAGGGGAGTATATATAAGTGTACA ACAATATTACTGTCTAACTGAACATTTCTTTAGGCAATTGGCATATTTCTTGAGAATATTCA CGGAGAAAATATATCCAATGGAAAGAAAGATACAGAGGAGAAAATGCAGGGAATT[T/A]CAG CATGAGAGTTAGTAATAGAACGAAGATTTTGCAGCTTCTCATGCAGTGCTGCTCTCTTTTGC TCTTTGGAACCATTTTGTGTGCTCTCTCTTCTCTCTCTCTCTCTCTTATGATCCTTCTTCTGGC TTTCTTTCTCTCTGGTGCTTTTCTTTTGCTATCAACTCTGCAGAGTTCTGTG
>Gm02_ 4789057 3	ATACAAAAGCTCATTCCAAGTATCTTGAAGGCCAGGTAAGCACCAATGCACACAATCAGCA TAACCTAACAGGGTTAGCTAATTGCTCTGGAGTCAATGGACTCCATTGCTTCTTGTAGATTGA AGTGTGTGCATCTCTACGATAATTGGATAGTTGTGTGATGTTCAAAAATGTAATGG[T/G]CCA CTTTAGATTTGCTAAACACTCCTCCAATTACTTCCATTATACTCTTCTTACAATCAGAGCCCC AATATGTGGGATCATCAATTAGAGTGGTCTCATTGTAACAATTCCTCCCGGTTACCTCCC CAGTCTATGCTCCTGTGCATATATTATTTATGAAACATTAGGTTATTCACACA
>Gm03_ 1380611	TTTTGCTTCACATTATTTAGGGTCTCCAATCACTTGCAACCATTTGCCGCCGGTAGCATATG CTGATATAAGTTACTTTTTGCTTAGACATATCCAAGACTTTGTTCAATTTCCCTTTTGTCTGA GGTGTGTGCATTCTTGTTTAACTCCTCTTTGTTCTCAGACATCCCAGAAAGGA[C/T]CAAA AAAAAAAAAAGAAAACCATAATCACAAGCAATAAAGGGGAGAAAACAAACATGTCATTAC AGGAACTCTTGATAAATGCACAGCATGTGATAAGACTGTCTATGTGGTTGACTTGTTAACTC TTGAAGGAATTACTTACCATAAAAACTGCTTCAAATGCAGCCACTGCAAGGG
>Gm03_ 1998320	ACACCCTACTACACAACGAAAATAAGTAAATTTCCGGTGATAGACATTTGTGTCAAACAAAA TAACACGAATGCCATGGGGGGTGCCAATAGCAACCCACAATAGGGGGCTAGAGAGAGAGAG GAGGCCCAAAAGGGAAGAAAGAAAGATAGTAATATGATTTTCAATTTGAGAGTATTTGA[G/T]T TGCGGCACAAACCATTCACTGTGTGCGAAAAAGAAAGAGAAAAAAGATGAGCGACGGA GGAAAACGACAAGGTGGAAGAGCGTTGCCGCCGCCCAAGAGGCCCTCCGCCTCCTCC CCCCTCTCGCCCTCGCCTCGAACCCGTCGATCGCGAAAAGGTACTCACTCTTCTTCTTCTC
>Gm03_ 3105779	ATCCATCAGGCCACATTCGATCCCGCCAGACACCTATTAATAAACAATAATATTAGAGCTGC CACATGTATCTATCAAAGTTATCTATTTAAGAGATATTATAAAGAAGTATACCGGCATTGATC ATAGGTTCAATTGTAAAGGTCTGTCCGGCCTTCATCACACCAACTGCTTTATTTT[C/C]CTGAC AGTGATTAAGGTGTATACATTCCATATGTTCTCGTTCTACTTTCTAGCAACCTTTGTTCTTTG TTAGGGGGTACGGGTAGAAAAGCTAGAGGTGCACGTTGGGAAAGAAAAGTTTGTTTAAAGA AAAAATGGCACATATGAATGAGGAAGCTGGCTGCTTCTAGCAAGTAATGGTA
>Gm03_ 4586070	TATAAAAAATATATTAATAATATTGAAAAATAAGTATTCATTATTTAAAAAATTTAATATTATT AAATGAATAGAATAAATAACAAACATCATTTTAAATATATGGATAAGAATTTTAAAGAATACGTA TTTAAAAATGAAATTATAACATATATATATATATATATATATATA[G/A]AGAGAGAGA GAGAGAGAGAGAGAGAGAGAGATAACGAAAAAGAGGTTGAAAAGATCGCGTATTATTACTT ATGCAAAAATTCACCCATTAGTCGGTTGGTAATAATTATCCTTTTTGTTGCACCTAATAAGA

	GTCAAAATAAGGTTGGCCTTCAATGTAAACCTCTCGTTGCAAGTTTAT
>Gm03_ 3589237 3	TGGCTCATGGAGTGTTAAGGACAACAAGGCCTTTGAAAAGGCTTTAGCTGTTTATGACAAG GACACTCCTGACCGTTGGTACAATGTTGCTCATGCTGTTGGTGGCAAACTCCAGAGGAAG TGAAAAGGCACTACGAACCTCCTTGTTCAAGGATGTTAAGCATATTGAGTCTGGACGTG[C/T]G CCATTCCCAAATTACAAGAAAACCTTTCAGGGTCAACTGATCAGGAGGAAAAAAGGTATC ATTCCCTTTATACGTTTGTGTTTGTGATCCGTATAAGAATAGAAGGTAGATATTAAGAAAA GTTTATAACAACGCACCTTACTTAACCATCTGAGTTAGATCTTTAATCTTATACA
>Gm03_ 3639987 4	CATCTAACTTCTTCAAAGGCTGAAGTGCATAAATTGATTTAACTTATAGAAGCTTTATTCAT TTTACTTTTTTATTCTTCTTCTCTTGAAGAAGTTTATCCAAACACGACCTAAATTATACTCCT AAATAGAATGCTTTTGGATGTTTCATTTCTGTTAAGAATAATTGATAGATGAG[A/G]TACCCTC CACTCAAGAAAAAGCGGCCAAGAGTAGCGATGACTAGCCGGGACCAGAAAGCCAGGGTGG ACGAAATACACAGTCTGAAGCCTGTCTTGAATCAGCAGGAAGTTCTTCATAGATCCACC TCAAGATGGTGATGCCAGGGGAATTATCCTCCTTCTGAACAGTGCTGTGCATG
>Gm03_ 4057250 2	TTCTTCTGGTCCTTAAGACCAGCATCCTTGGGGTTGGGAACGTTTCTGGCACTTGTGGCCA CAGCCAGAGCAAGAACCAACACCAAAAACACACATCTTAGCCATGTCTCTCCACACACTAG GAATGTACAACGAGTTATTGCACTCCTTTTGGTTACTCTACTATGAGTGTGTGC[G/C]GA AGTGTTAAGAGGAGCATGTTTTTATAGGGGGGGGAAATAAGCTAACCGGTGAGGGGACA GTTGAGGATGGTAAATGAAACATTAATGTGACTGCAGAGAGCCTTGCTGTCTATGTCCCAA TACAATTTAAGACCAATTCGTACACTGCAGGCGCATACGTTTATTATCGGTGTGTA
>Gm03_ 4206283 1	ACTGTCAGCAGGACAAGGGGGTGTCAATCACATTGATGGACACCTAACAGAAAATAAGATG CAAATGAAACAAAAAGATCTAACCTCAAAGCTCCACTGAAGCTCCATTTTCAGCTGATTTGT CTTGAGTCGGGGGATCACAACAGAGAGAACTGGCATGGTTGGCCTATATTTTGCTA[G/T]TA ATCTGAAAAGTGCATGCGATGCTTTTAAAAAGCCTTACAGACACAAGTGGAAATGCAGGG AAGAAAGATTGATTTGGACAACAATTGATGCATAATTCATTTTCCAAGATACTAAATATGA ACATGTTTCATATCAAATTATTCGTTGAGACACTACCTTGACGCCCTTCCAGAAG
>Gm03_ 4386579 7	TGGTGTGGCAGTGGAGTTCTTAATCCGATGGTGTGTGTTGATAGGGACTAGGGAGAAAGGA ATGGCCACGTGCTGTTAGGGACTGCTTCCACATCTCCGCGAGGCCAAACCCAGTGTCTAC TCTCCACCAGAGAATTCCACGCCCTCTCCTTTAAACCATAATGCACCCCTTGCCCTTTCT[A] CCACCACACTCGCTTCTCTTGCTTCCACTTCCATTCCACCATGCCTTCTAATGATCTCAAAC ATCGGGTCACCACCACTACAGCAACCACCAACGACGTCAATAGGACCAGCAGAGGCAAAC GAATGCTGATGGCCAAACGCGGCCCTTAAGTCACTCGCCATAGCCGTTGTCTCCTCT
>Gm03_ 4622405 7	TCAGATCCATAAGAGAAAAAATAGATACACAACCCAAGGAGAGAATGCAACTTGAAAAGCA GCAATGGCTCGTATAACTTCATATTCAAGGCTCGAAGACCATGCTCTCTGCTTCTCGCTTC ACGGATTCAAAGAGCACAGATGATAAATAACGCTCTCCAAAGCTTGGGAAGACCGC[T/G]TA CTCAGGTTTTTCAGTGGGCTTATCAGTGATGGAATATATATAGTAAATATAATAAGAACAAAGT GCATCTAACTATAACAACCAAAAAAATCCACATAACTAATGCTAGAAAGTTTCTGCCAATGAT CAGCACTTAAAGTGCAACGATTCTGAAAAATGTTGAGCACATTGATACTGATAT
>Gm03_ 4716319 4	CCAAGAGACAAATCCGAACATTCTTCTCACAGTCGATGATAATAAAAAACAGTATAGCTAA ATTAGAGACCATGAGAAGCTTATAAAGCAAGAAACAGACCTTTTTAGCATTCAGGAAATAA CTCTGTTCTCATACTGCCATTAGCCTAAAAACAGGTAATTTGAAAAATAATTAGAG[C/G]CTTA CTTGCACTCTGTTACTATAAGGTTAATGGCTTTGTTTCTTTGTTTTCTTTCCCAACAGCAC TGCTGGTGATGCTTCCAGAACAACCTTAGGCAAAATCTAAATGCCATGAAAAAATGGATG GTTTGATGAGGTCAGTAGTTACACAAAAGAAAAGCATAACAATAAACCTAAGA
>Gm04_ 739976	AACCCAGAACCTGCTGTGCCGGTAGCGGTGGTAGCGACACCAGTGAGTTGGACGGGATTA CCGCGCTCGTCAGTTAAGATAACCGGATTACCGTATTGATCGGTGAGTGGGACAGGGTTG CCATGCTGGTCTCGTAGTTGTGCTTCAGCCATTTCTTCAACTCTTGAATAGTATCTGC[C/T] GCTGAGTGCTGAGTACTTAGTTATATATAGCTCAGTCTGGGTAATTAGTTCCTTCCTCTC AATGGCGGTGGAGACATGCATGTAATTCGGCGGCAACACGTGACAGGAGAACACACAGAC ACCTCACACCCGTTTGCCACCTAACAGGACACGTGTCTCTTTGTTGTAGTGTACAC
>Gm04_ 4586273	GTATTTGGAAGTATGAAAAAATGGCTCTGGTGATGCTGCTATCGGCTCAAGTAGGACTTC AATATGTTCTGGCTAGTGATTACAATGGTGACAGGGGAGAGCACACTTTGTTGATCGGTTA CAGAACGTTGGTTCAGCTCGGGCAGCTCCTGTTTGGCTTGAAACCAGTGATGCAGTT[G/T] GTAAATACTTATGTTTATATGCTTCAATTGTGGGATGATTCTCTGTCCATTATTTTCTTTAAC AATTTTGCATGGCAGCAAATATGTTTTAGAGCTGCGGGAAGAGGCTCGCGATCATGCATG CTTACGTAATGCATATTTTGTAGCAGGTGTGTCTTCAATTCATGCTTTGATGAAGT
>Gm04_ 4586274	TATTTGGAAGTATGAAAAAATGGCTCTGGTGATGCTGCTATCGGCTCAAGTAGGACTTCA AATGTTCTGGCTAGTGATTACAATGGTGGACAGGGGAGAGCACACTTTGTTGATCGGTTAC

	AGAACGTTGGTTCAGCTCGGGCAGCTCCTGTTTGGCTTGAAACCAAGTGATGCAGTTG[T/C]T AAATACTTATGTTTATATGCTTCAATTGTGGGATGATTCTCTGTCCATTATTTCTTTAACAA TTTTGCATGGCAGCAAATATGTTTTAGAGCTGCGGGAAGAGGCTCGCGATCATGCTGCT TACGTAATGCATATTTTGAGCAGGTGTGTCTTCAATTCATGCTTTGATGAAGTC
>Gm04_5096967	TAAACATAAAGTCTATTATATATATCAATTTTGTGTTTTTGCATGTTGATTTTCAATCTCCTTT GCTATATTTTACGTTATTTGTAATAGTTAGTTTTGGGTGTTGGTAGTTTACTAGTTATAACTT TTGTGTATTCTTTTTGTTGATATATCCCTTAAACATAATTCTTTCTATACAA[G/A]ACATGGGT ATGGAAGGTGGCAAGCTATTGTTGATGATAAGGATCTGAAGATTCAGGAGGTCATATGTCA GGAGTTGAATCTTCCCTTTATAAACTTACCAGTTCCAGGACACGTTAGTTCTCAGGCACAAA ATGGTGCAAATTTGACAAATGCAGAAGTACCCAACAGTCAATCCAAGGA
>Gm04_5581132	TGTTTCTTGTCTACTGATGTCGATGTTCTCAACAATGTAGGGAATTAATTTTGTTAACTTTT TTTTAATGTTATACCCATTTGCATTTCTAATTTGTGTTTGATGGATTTACAGGTTCACTCTA GGTTTGAGGTGGATATAAAGCAGCTTCTGAGCAGATTGATACCTCTACCTAT[C/T]GTACG TATTTTTTCATCCATTAATACCTCTACCTATAGTTTCATATAGAGTTTAATTAGTTGACTGAAAA TTATAATTTAGAATTTTAATTTTGTTCCTTCCAAATTCTTCCAAATTGAAATTGAATTAGAATA AATTCTATGTTTGTATTGTGTGCTTAGGACATATAATTGAATTTTAA
>Gm04_6375455	GCTTCTGTGTCATGTTTTTGTTCATACCAACTGTGGCATTTCCTGCATCATATAGGTAAGT GGCTGGGAAATAGGCAAATACGATGCAACTGGGCAACTAAGGGTGCTGGTGGCAGTTCCA ATGAAGAGAAGATTAATGACAGTCAAAATGCTGTTGTGCTTACAAATGGATCTTCA[A/G]GT ACCATGTACCTTTCTAGATAGCTAAGAGCCTGAGCCTACATTTGTTAAACCATGAAAATATA GATGAAGGTTCTAGTTTAAATTTGTTTTCTTCTCCAGATCTGGGATATTTTATTGGCTGATAT TTGGCATTCAATTATGTTGAATGATATCATACTGATTCAAATGTATAATGGAG
>Gm04_7079456	CTAGACATATCATATTTCAAATGCTAATTAATATACAAGGCTTATAAGGAATGGAAGCTAG GATGTTTCATATAAAGTTAATGGATAGTTTAAATTTTCATGTTAAATCTTTATCGATCTGCCTC TCTGTTATATTTTGATAGATATGCATGTGCTGAAATATAATTGTTTTGCGGCTG[G/C]GTAGA TAATGGAGAAAGCTGACTCCAGCAGGAGAGCGTCCAAAAATCAGCCAAATGCTCTTCTCTG GCCTTGTCATCCTTGCAAAGGATGTAGGTAGCCTGGCAGTCAATGAAAAGCAAGTGTTTCAG TCCAATACTTAAGAGATGGCATCCTTTGGCAGCAGGTCTTGCTGTGGCCACCCT
>Gm04_7891337	AGATTTATATCATATTGGAGTTCATCACGGGTGGTGAATTGTTTGATAAAATTTGTGAGTAGC TCAAGATGGAACCTTTTTTTTTTTTTTGTAGATTTTACTCTGGCATTATTAATCACCATTCAAG ATATTGGAATGTTTGATGGTGAATAATATGATTGTGCAGATACACCATGGCC[G/T]TCTTAG TGAAACTGACTCTAGAAGATATTTCCAGCAGCTTATTGACGGTGTAGATTATTGCCACAGTA AGGGAGTTTATCACAGAGATTTAAAGGTTTGCATTGCGTGAGATCTGAAAAGTTGTCTATTG TCCTTATGGTGATTCTTCTAATTTTCTTTTTTGCCTTGTGCAGCCTGAA
>Gm04_8124418	TCCAAGTCCCATTATTGGGTCTATAATCGGCCAGTAGCAAAAGAAACATATTCGGGCAAG GCTTTTGTATTTTCACTCCCCTTGTACATGTGGTGCATTTGGTGGCAATAGGGGGCTCC GCCAGTGCCCTCCCGAAAAGGGAATATATTCTTGGACCACATGCATTTGTGTGACC[T/A]A GAGAAGATAGAATATCTCAATTGTCTAAAAACTTTTGGGCACCAGCATGAAAAATGCAGGC ACTTCTAAAAAAGGGAATATTTTCACTTACATGTGCGCATGGAGAAGAACTTGATGGCAAG CAAGCCCTGGTTGAACCTTGGTTTTAAGGAAAGTAATGTACAACTTCTGGAGGTAAA
>Gm04_8124424	TCCCATTTATTGGGTCTATAATCGGCCAGTAGCAAAAGAAACATATTCGGGCAAGGCTTTTG CTATTTTCACTCCCCTTGTACATGTGGTGCATTTGGTGGCAATAGGGGGCTCCGCCAGT GCCTCCCGAAAAGGGAATATATTCTTGGACCACATGCATTTGTGTGACCTAGAGA[T/G]GA TAGAATATCTCAATTGTCTAAAACTTTTGGGCACCAGCATGAAAAATGCAGGCACTTCTAA AAAAAAAAAAAAAAAAACCCTACAATGTGCGCATGGAGAAGAACTTGATGGCAAGCAAGCCCT GGTTGAACCTTGGTTTTAAGGAAAGTAATGTACAACTTCTGGAGGTAAAATTACT
>Gm04_14625604	CCTTCCATCTATGTTCTTTTGGTGCGGAAAAATAAAGGATGTAGTTGTTATTGTCAAAATTC CTATAATATATTTTCAACTTTTTCTCTTGGGCTTTATTTTATATCTTAACTTAAAAATATCCA ATGTATAAAGTTTCTAATAGTTTGGCTTTGCACAAATTTCTACTGAAGTACA[C/G]AAAGAC GCAGAGCTGAATTAATCGGCATTATCACATCACAGCGACACAGCCCTGATTTATCTTTTTTC TGCTTTTGGTAAGTTACTCAAAGTTAATTTGTGAGTGTGAGATGATGCATGATTGCATGA ATATATTGTGGACATGTTCTTAAGCATTATCTGTTGTTTTGTTGAGTTT
>Gm04_14966203	GCTCCACTGCCTCCACCTCCAGAATCTGTACGAACAGGTTTGGTACCACCACCTCCTCCAG GTCCCGTGCGAATAGGTTCTGCCCCGCTCCACCTCCACCTCCACCTCCACCTCCACCTC CACCTCCTCCGGGTCTGTACGAACAGGTTCTGTCCCGCCACCACCTCCTCCTCCTCC[A/G] JGGTCATGTACGAACAGGTTTCAAGCCCCACCACCACCTCCACCTCCTCAGGTCCTGTACGG ACAGGTTCAAGCCCCACCACCACCTCCACCTCCTCCAGGTCCTGTACAAACAGGTTCAAGCC

	CCACCACCTCCTCCTCCTCCTCCAAGTATACGAACAGGTTTCAGCTCCACCACCTCCTCCT
>Gm04_44170876	CCAACAACTTAACAGTAACACACAACAGTGGATGCAAATTACAAAAAAGGAGCTCACCGGAAGTAGTGACAAATAAACCGATAACAATATCGAACCCAGAAATATAAAAGGGGAAAGAAAGAAAGAAAGATAAAAAAGACGAATTAAGTATTAAGAAAAAGAAAGAGAGAGAG[A/G]CCTAAAGTAGCTGAAAAGAGAAAAGAGAGAGATGAATGAATTATATAGTGAGCGAAGAAGAGAAGAGGAAGGAATAAAATTGTGGTGAGGAATAGGCGCTAACCTTCCCCTGCCCTCCTTTCC AAGATATCTCTATCTCTAATCGATAACTCAATCTTAAATATCTACCGTAACATTTT
>Gm04_46190568	GTTGAAGAACATATGCAACAAAAATGAATAGCAAATCAGGAAAGCTACTCTTATCTTGCATAACACCACCTAATCTCAACAATCACAACATAAAAGAAAAGAACCCATTATCCACTACAACTTTATTTATTTATTTGTTTGTGTTTGTGTAATCCTCAGGCTATAAAACGAGTTAAGAC[C/G]TACGTGACAGGATCTGAAAGAATTGCCCTGAGTGACTCCCTGACTCGCTCGAATACATAGCATCCTTCTCCCACTAAACCCACAACCACTATCATCCACCATCACTTTCTGCCCAGAAGCCCCCTTGCCGCCGTGAAGATAGTGCGAATAGTACTCCTCACAGTCAGTTTCAGCAACG
>Gm04_46190569	TTGAAGAACATATGCAACAAAAATGAATAGCAAATCAGGAAAGCTACTCTTATCTTGCATAACACCACCTAATCTCAACAATCACAACATAAAAGAAAAGAACCCATTATCCACTACAACTTTATTTATTTATTTGTTTGTGTTTGTGTAATCCTCAGGCTATAAAACGAGTTAAGAC[C/G]TACGTGACAGGATCTGAAAGAATTGCCCTGAGTGACTCCCTGACTCGCTCGAATACATAGCATCCTTCTCCCACTAAACCCACAACCACTATCATCCACCATCACTTTCTGCCCAGAAGCCCCCTTGCCGCCGTGAAGATAGTGCGAATAGTACTCCTCACAGTCAGTTTCAGCAACGG
>Gm05_806881	AGTAAGCAGAATAGTTTCTCTTTATCCAATTTCAATTCAACTCACCCAAATATGACAGCGTTCACAGCATTATGTTGTTATCCTGAGGTGCCCGCTGATTCTGCAGGCGGGTCTTGTTGCAGCCTCTTGAAATCTCTCATGAGCCTCTTCTAGCAGGAGTCGACATTGATATCAC[C/T]TAAAGAAAACCAACAGCCACAAAGAGAAAACGGAAATAACTTCAATTTTCTATTAATAAACTATTAGAAGCATAACCAACAGGTCAATTCAGATTCCAACACAGACTAAAGACGTGGAAAATGAGGGGTAACACTAAGAACACATTTGAAGATATAATTCAAGAATTTAACTTGCAAAAT
>Gm05_1305399	CAGATAATTCATTCCTTCAATCAAGTTCTTTTTGTTAACGTTAGTTTGTATCCACTTAAAGATGAGAGTGACGATTTTTCTTTTTCTTTGTTATTGTTATTTTGTCTTGTACAGGTTTCATCTGATGTTAACATAGAAAAGGAGCAAGATGAAGAATTTTCGCTTTCCTGTTGACCA[G/T]GCATGTATTTATTTTTTCTTGCCAAAAATATTAGTCAAAATATTTCTTAAATTCTGTGATTTTCTTATGTGCTTGATGTAACCGTATATGTCATAATATCCTTACTGGAAATATGAAATTCTCTTCAGCTTATCGTGAGAGGCGGGAACAGAAGGCTAATGCAGGACATAGGTAAGC
>Gm05_1964837	GTGGTGATGCCACCGGCGTTACCCCGACCACCGGGATGCTTGCGGTGCTTCCCGATACGACCGTGACCGGCGCTGACGTGGCCTCTCTTCTCCTGTTCTTCTTGAACCTCGTCGTCATTCTCTCCTCCGCTCCTCTCTCTACACACTATCTCTTCTCTCACAGAGTGGCTG[T/C]JACAAACGAATTAGGGTTTTGGGACTTATAAGCCTGTCCACTGCTCGCGATTTTACTTATTTGCCCTCTCTCACAGTCACACATCATGTTGGTGTTTTTTATTAATTAATAATTTACTAGAGTAAATAATATCTTTAAATATAAATTATATTATAATTTAAATTGTATTTAATACATA
>Gm05_2011564	GATAGTAGGTTTATTTTGGATGTTAATGTGATAGATGGTACAATGATGGCTCCATCTACTGCATTTACAAAGATCTGGCGGATGCGCAATAATGGCACTATAGTCTGGCACAAGGGAACCTCAAATTGTTTGGATTGGAGGAGACAAGTTTAGTGACTCTCATTAGTTGACTTAGAGGT[G/A]TGATAAATATTCATATTGTTGTTTAATTATTTCTTTTAAAGTTTATTAGCTTTTCTCTGGGTTTGT TTTGCTTGTTATAGAATCTGTGGCATGCTAGTTTTTGAGTGCTCTCTCTTGTAAACATCCCA GGTTCTGAGGATGGTGTCCCTTTGGAGAAGGAACTTGACATTGCCGTTGA
>Gm05_2011565	ATAGTAGGTTTATTTTGGATGTTAATGTGATAGATGGTACAATGATGGCTCCATCTACTGCA TTTACAAAGATCTGGCGGATGCGCAATAATGGCACTATAGTCTGGCACAAGGGAACCTCAAC TTGTTTGGATTGGAGGAGACAAGTTTAGTGACTCTCATTAGTTGACTTAGAGGTT[C/T]GAT AAATATTCATATTGTTGTTTAATTATTTCTTTTAAAGTTTATTAGCTTTTCTCTGGGTTTGT TGGCTTGTTATAGAATCTGTGGCATGCTAGTTTTTGAGTGCTCTCTCTTGTAAACATCCCA GTTCTGAGGATGGTGTCCCTTTGGAGAAGGAACTTGACATTGCCGTTGAC
>Gm05_3005246	AATGTTAGTTTTGAAAAATAGGAAAGAAAGAAATGAATTCGATGACAGAGATTGAAAGAATG AATACAAACCCGCGCTGTGCATACGACCCATGGCTGGTGGCTGTCACTCGCTTGAAGCT GCGCTTCTTCTGATTAGGGTTGCGACTACGACGCTCTTTTGCACACTCCACCACTCT[A/G]JAAACCTAGATTATGCAGTTTATGTATTTTATAAGGATAAAAAGTGCAAGCCCAATAGGTTT GATAGTTTTTGAAGGAAATTGCTACGGGCACCCGGCAAGATTGCTGGTGCACCCAGCAAT TTTTATAATTCCCAAAATACTCTTAAATGTTTTTACGGAAAACTTTCTTTTACA
>Gm05_3461049	AATTTTTTTTTAAGCAAAATTAATTTATACAAACCCATTAAAAAATACTTATTTTATTAATAA AATATTTTCCAAGAAATGAATTTAAACAAATTCACCTCTTATTAATGGGATTGGGAATTGA

5	TTCTATGACGAACGCCGAGAATTTTGGTCTCTACCGTCTTGACTCTTTAGG[G/C]TCATAATT AGATACAATATTTCTACAATTCTACACATTTTACGTCCGACAATGTAAATTAGATTTACATGA GATTGATATCAACTTATTTGTCCTCTGTATGAAGAGGTTGCGAATACTTGTGCATAGAGGAAT TAACTTTTAAAATTTAAAAAGTCAAATGAAGTGCCAGTGACTTCATT
>Gm05_ 3461049 6	ATTTTTTTTTAAGCAAAATTAATTTATACAAACCCATTAAAAAATACTTATTTTATTAATAA ATATTTTTCCAAGAAATGAATTTAAACAAATTCACCTCTTATTAATGGGATTGGGAATTGAT TCTATGACGAACGCCGAGAATTTTGGTCTCTACCGTCTTGACTCTTTAGG[T/G]CATAATTA GATACAATATTTCTACAATTCTACACATTTTACGTCCGACAATGTAAATTAGATTTACATGAG ATTGATATCAACTTATTTGTCCTCTGTATGAAGAGGTTGCGAATACTTGTGCATAGAGGAATT AACTTTTAAAATTTAAAAAGTCAAATGAAGTGCCAGTGACTTCATT
>Gm05_ 3679675 3	AGCATCAGATTTAGATGATGTAGAATCAAGGCCAAGACATTTGAGGAATGCTTCTTTGGCTA AGTTGAAATAATAGCAAGGGTGGAGGAAATACACAGACGAACCTGAAGCCTCCCCTTCTGC AACTCTTCTCTCACTATTCCTATCTCACCTTTAATACTTGTGTTCTCTACTCCCT[T/A]CCAT GTCAATTTACTTTTAGTCAAAAGGAAAATAAATTATAATGCATATTTTCATGTAAATTTTGT ATGATTAGATTATAGTTAGAGCCTATGATCTAATAATATTTAAACTCGAGATTCAATGTACAC AAAAAGATTCCCATCTTTCTTTTATGCTTTAATAATACTTATGAGAG
>Gm05_ 3727103 2	TTGGGAATTCCTATTTTAAGGTGTTGAGAAGTAGTGTATTGATAGTTGGGAGAAGGGTTA GGTCTTGAAGGTGGGATTTTCATGTGGAAGGTGGTTTGTTCAAATCGAAGTGTGAGATAT GCTGAGCTCTGATGATGGAGACATGGATGTGTTCTTTGACTCAGTGGATAGTTTGT[C/G]A CTCAAGATTCTGTTTTAGCCAAAGAAGAGTTTGGTCTGGAAGGTGTGGTTATGACGAAATT TGGGTGAAGGAGCCTGTTAGTGTGAAGGAGAGGAGGGAACGTTTTCTGCAGGGTTTGGGC TTAGCTGATTCTTCTTCCAAGGTTTGTTCACAAGAGAAAATGAACCTAGATGACTCA
>Gm05_ 3866506 1	CTCTTGCAACAAGTGCAGGCCTCACTCCCGCGACAAAATCTTCATCCTCCCTTTTGATCACT CCTCTCCTCCTCTCACAAACACTTCTCTTCTCTCAAGCCCCAACGGGATTTTCAGGTC CCTTGTTTCCAAGCTCTCCCGCAAGAGCCCCATGTCTTCTTCTCCTACATGACCTC[C/G]TC CTCCTCCTCCACCTCCCGAGAGGAACATTGGAATTTGGCTGTGGCCGAACCTCTCCACACA AGCTTCTCCACGCCACCAGGAAAAGAGACGAGGCCCTCCTAGAAGCTTCCAGACTCATGC ACTCCATGTCCAGTTAGAGAAGAAGCTCAACAAGCTCGAACTCTATTGCCACACCTT
>Gm05_ 3941063 1	AATTCAAGTGAAACCGGTTTTAAATTGCTCCAAGCTGCAGCCTGCAACACTTCACGCCCTA CTTCAATTGAACCAGTAAACTGACCTGAAGATCAACACAATCAAGGAAATTGAAATTTTAC AAGTCAAATCAATTAAGAGACTAGTACTTGTAAATATTTGACATGTAACCTCTTAT[A/G]ACC GCATCTATGTCCATGTGTGAGCTTATTGCTGCGCCAGCAGTTGGGCCAAATCCAGGTACTA TATTAAGCACACCATCTGGGATTCCAGCCTTTTAAAGGGACAAACCCAATAGAATTAGCCT AAATTCAGTAACTTTGTGAGGGTCATCTTTGATAAGGAAATGAACTATGTC
>Gm05_ 4140548 4	TTAAAAATAAAGAGCTATAATATAAGATTTTAAATTGAAGAGAAAACGAAATTTCTAACA GATTTAATTATTAATTAAGTAACTGAACTCGAGTGAGGGAGGGTTTCATTTCTCGGATAGGAAGT GAAGTGAGAGTGACGAGGAGAGGCCACCAAGCAGAACTGCTTCAGGTTGCCAACT[A/G]GTA AGTTTCGGATTCTCTTAAAAAGCAGTATTTTCTTATCTGTTGTAATTTGACTTCGGATTAACC TTCTGCCAGTTGACTTTACAACTTGGGATACTCCACATGTCATAAACATTCAACGTTGTAC CACTGTTTTTCTTTCATGTTTCTAGTAATTTACCCACACTTTATTTGCCCC
>Gm05_ 4187381 5	TTGACCTGATATTAACCAACACATCAGTCATATAGAAAGAACAGAAAAACATGAACCTAAA AACAAGATCTGTAAACAGTAATTAATAATTACCCCGAAAAATATGCCTGAATCGCCACTCGT TCCCGTGCGATCCTTGGCCACCAACTCCTGGGTAGGTGGCTGCTTCGTATGTC[C/G]TG TCAACAAACACGAAACAAAGGAAGACTCCCCCAACTTCAGCACAAGGAAACAAACAAAGTA CGTGGAAGGAGAAGCTAAAAGGGTAAAAAAGAGCAGCAAAACCAACCAGCGGAGGGAG GCATTCATCGGCATGTGCGCTCAGCACCGAAAACCCACCGTGAGTGCTCGTATCAGA
>Gm06_ 2019325	CAAACTAAACCAAGGAAGGATCTACAGATTCTATTAATTAGTAGCAGCATAACAGTATCG ATAATTCTTAGGACTGTGCTGCTTATCAACATGAAGATTGAAAATCATAACAATAAAAGAGC AACTTTTAGGAGCAAGATGTTTAAAGTGGAAGAAGAAAATAAAATAAATGGAAGT[C/T]ACC GAAGAATGATCCAAGCTGTACATTACAATTTTGTATCTGCAGCTCCTGAGATACCCCTTT GCAAGACCCATCATTACAGACGCTCAGAACTACAAAACCTGATCAGTTATTGACTACCATAAA TTTTATTAATAAATAAAGAAATCTTAGGAAGCTAAATCCACACCTGGCTCAG
>Gm06_ 8655798	CGTAAACGGCTTCTACTCGTTCCTGACGCGAGGAATCGACGATCTAGAGAGCGTGTTCCCTC AGCACAAACTTCATGTCAATTCATTCCTCCAAAGAGCACTCTCTTCTAAGATCATTCCA CACCCAATTGACTCTCCTGGTTTCAGAACTCCACCTCCCCGTGCGCGACAAATGGC[G/T]C GACGAGTACATGGACGAAAGCTCCAACTCTGGGACGCTGCCACGTCATCAAGTCCGGC ATTTCCAGCATCGACACCTACTACTCCGCCGCCCTCAACATCACCTCCTCCCTCGACTCCC

	ACCGCCCCATCACCCCCCAACTCTCTCGCCAGGTCACCAATTTTTCTGTTCTCTCTATT
>Gm06_865844	GAGAGCGTGTTCCCTCAGCACAACTTCATGTCAATTCATTCCCAAAGAGCACTCTCTCTCTAAGATCATTCCACACCCAATTGACTCTCCTGGTTCAGAACTCCACCTCCCCGTCGGC GACAAATGGCTCGACGAGTACATGGACGAAAGCTCCAACTCTGGGACGCCTGCCA[T/C]GTCATCAAGTCCGGCATTTCAGCATCGACACCTACTACTCCGCCGCCCTCAACATCACCTCTCCCTCGACTCCACCGCCCCATCACCCCCCAACTCTCTCGCCAGGTCACCAATTTTTCTGTTCTCTCTATTTGTTTAACTGTATCCTCCTTCGAAACAAAACAAAAAAGATTCT
>Gm06_10814320	CCCGAGAGTAATAAGGCACTAGAAGAAAACCTACATGTCAGCAAAGTTCATCTTCCCAAAATGAAATTTGTAGGATGGGCTGGGTAATTTGGACCTGATATTGTTTTTCATAGATAACCATGATATTCGTAGACCAGAAAAATGAATATACAATGTCATTAAACAGGGAATAAAAAAGTT[G/A]CTCTTACAATTTCACTTCTGGTCATAACATGAGGAACCTACAATGAAGTAGGGGAAACAATGACAAATTTCAAATGGTTGAAGGTAAGAACTTCCTGTACAAGATGACAATCACAAAGCATGAATAGATGGGGAGCAGAAAGGAATGCTACCAGTTCTCCTTAGCAATTTTTGACTTTTC
>Gm06_10888465	TGATGGCCATACCAATAGCAAGAAAACATGAAGCAGCATCAAAGGGGGCCACAGGTCCAAAGACCAATGTATCAACAAGTACATTTCCAAACAGCCCAGACAAAATAGCAACAAGACCATTTCGAAGAAATATAGCCTTTGAGAATGTTAATGATAGCCATTGTTGATCAAAGCCCCTC[C/T]GACGATGTTTTAAAAAACAGGCAATTCAGAATCATAAATACACAACGTAACAAGAAGGAAAAACAAAGACCAAAAAAAATTTGTAACAAGAAGTATACACTGAAAAATGTGTCTCAGCTCTACACTGTGGATTCAATATCCATTTAACGAATCCAATAAATACCAAAATGCAACTTTCTCAA
>Gm06_10888466	GATGGCCATACCAATAGCAAGAAAACATGAAGCAGCATCAAAGGGGGCCACAGGTCCAAGAGCCAATGTATCAACAAGTACATTTCCAAACAGCCCAGACAAAATAGCAACAAGACCATTTCGAAGAAATATAGCCTTTGAGAATGTTAATGATAGCCATTGTTGATCAAAGCCCCTC[T/A]GACGATGTTTTAAAAAACAGGCAATTCAGAATCATAAATACACAACGTAACAAGAAGGAAAAACAAAGACCAAAAAAAATTTGTAACAAGAAGTATACACTGAAAAATGTGTCTCAGCTCTACACTGTGATTCAATATCCATTTAACGAATCCAATAAATACCAAAATGCAACTTTCTCAAT
>Gm06_45353845	CCTTGATAAACCTGAACCTCAAGGGAAAAAAACACACTTTGTTAAGATCCTCTCTCATTTGGCAACGATTTCCGTATCCCAAAGTACCAAACTCCAAAAGGTGTCAACCTGCCACTCCACACGCACATGGCAGCTACCAGTAAGAGGGAAAAACATTAAAAAACAAAGAACGTCAGC[G/C]CATCGTTGTAGTTATCACTATCATCATCATATTACAATTCTCCACTCCTTCCCTAAGCTGCCGAATCAGCACTTCGTCATCGATTCCAATACGGCGCCGTTTTCTTCGCCATCAAGCGCAACGCATAACAATCGTCAGCGCCACCATTTCCTTCGGTTCCCAAGAATCAATGG
>Gm07_2019872	CAATGATGCTCCTGCCCTAAAAAAGGCAGACATTGGGATTGCAGTGGCTGATGCAACCGATGCAGCTCGAGGTGCATCAGACATAGTTCTTACAGAGCCTGGTCTGAGTGTGATTGTGAGTGCAGTTTTGACAAGCAGAGCCATTTCCAGAGAATGAAGAAGTACACCATCTATGCTG[C/T]TCTATCACAATCCGAATCGTGCTGGGCTTTATGCTGCTAGCTCTTATCTGGAAATTTGATTTCTACCTTTTATGGTTTTGATCATTGCAATACTAAATGACGGAACAATTATGACCATTTCGAAGACAGAGTGAAGCCATCACCTCTGCCAGATTGGAAGTTGAATGAGATTTT
>Gm07_2482464	TCCATAATTGACAACTTAGAGTAAAGAACGAAATAAATACGTTAAAAATACAAATGATTAAATGAAAAAATCCACATCACAAAATGTGCGAAAAAAACAAGAAGGGTTAGTTCCAAGATTTGATACCCATTCCCTAGTGCTGAGAGCTCAACTTCATTATGCTGTTGCATATACCTC[G/T]GCGAAAAATAAAAAATAAATAAATAAAAAACAAATTAACAAAAAAATTAACAAAAATCAACCGTCTGCGCTACAAAATTAACATGTCCGAATAACTGAAAAATTTACGAGATTGATATCGAAGGGAAATGGAGTGAGTACCTTTGGCGAGATTGACGTAGAGAAGAGGGGCTTTTTGGT
>Gm07_2482465	CCATAATTGACAACTTAGAGTAAAGAACGAAATAAATACGTTAAAAATACAAATGATTAAATGGAAAAAATCCACATCACAAAATGTGCGAAAAAAACAAGAAGGGTTAGTTCCAAGATTTGATACCCATTCCCTAGTGCTGAGAGCTCAACTTCATTATGCTGTTGCATATACCTC[T/C]CGAAAATAAAAAATAAATAAATAAAAAACAAATTAACAAAAAAATTAACAAAAATCAACCGTCTGCGCTACAAAATTAACATGTCCGAATAACTGAAAAATTTACGAGATTGATATCGAAGGGAAATGGAGTGAGTACCTTTGGCGAGATTGACGTAGAGAAGAGGGGCTTTTTGGT
>Gm07_5667815	GCATAATTTTTATAAATTGTGTAGGCACTGACTCCAACATTAGAGTTAGAGGAGATTGATTACCAAAAAAGAAAAATGGTTAGAGGAGATCAACGAATTAAGTGGCTTGGTGCTACCGAAAAA AAAAAACAAAGTGGCCCAATTTTATTATATTTGTTTGTGTGTGAATAAAAAACA[G/A]ATTGTGCGTGTGGTCTACACAGTTAGTGGTGAATACTGTTCTATTCTATTTATTATGTAATGTGTGTGATTATAAGTCTTAACGACCCTGGAGAATAAAAAATGAAGTGATGAAGTGAAGTTTGTAACTAACTTTGACTCTATTGTTTTATGACAAAATAAATAAAAAATGTG
>Gm07_5855116	GTGAAAAGCATATTTGACGTGAAAACAACTACGGGATTGGGAGCAGCAGCCTTCCCTAGACCTCTCCCTTTTTCCCTCAAGAAATTGACTTTAGAGAGGCTGCCAAAACCTGGAGAATGTCTG



	GAATGAAGATCCTCATGGAATTCTAACCATGCAACTTCTACAACATGTTATTGTTGA[C/T]AA ATGTAAATGCCTTACAAGTGTGTTTCCCGCATCAGTAGCCAAAGATCTTGAATACTAGTTG TCAAAGACTGTGAGGAATTGATGGAAATTGTTGCAGAGGATAATGCAGATCCAAGAGAAGA CAATCTAGAGCTTACGTTCCCTTGTCCTGTGTGAGGTCATTGAACTACAAGGT
>Gm07_ 1195097 7	CAGATCCGCGAAACCATCGAAATGCTCTCAATAACCGAAAAAACCAATATTGCGACGCGG AAAAAGGGGGCAGAATGGTAACAGACGCAAAATTTTCAGGGAACAGAGGCTGAAGAGAGT GGGAGTGTGGGTAGGGTTTTGTTTTGTGAGAGAGAAAAAAATTAAGAAAAAATAAA[A/G] GAAAAAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG TAACCAGAAACAGAAATTTTAGGAGTGAGAAATATTTTTAGAGAGAGAAAGAGGTAACAA ACTGAGCAGTGAGAGAAGGGTCAAAAGAAGAAATAGGTTTTTCAGGAGATCGTAGGGTT
>Gm07_ 1720248 9	TCTCCCTGGTGGTTCTGCTCAGACGAAGGAGTATTATCCGACTCTTCGCGCAGAAGGAAG AGTCTTGATCGGTCTAGTTCTATTAGGAGAACTGCTGCTGCTGTTGTTGCTGAAATGGATG AGTTGAAGGCTGTTGCAATGCTAGTGCTAATGCTAATGCTAATGCGAGGGTTACTC[G/C]T GCTGCTTCTGCTGATTATGTGCATAATAATATTAATAATAATCCTTTGAGGGACTCCAATTTG AATTCGAATTCGAATTCAAATTTGTTGAGGGATGATGAGAAGGGGAGTTCTAAGAAGTCAT CGAGCCCCGGTGGAGGGCGTGGAGTATATGGGGATTGATACACCGGAGGGGAAGTA
>Gm07_ 3646081 4	AAAGAAATGAGACTGAAGATTATTGAGGAAGCTGAGGAGTATAAGGTGGCTTTCTATGAGA AAAGGAAGCTTAATGTTGAGACTAACAAGGTGCAAAACAGAGAAAGGGAAAAGGTAAAGTT TACACACAACATATATGTATCTTCTATTTTAATTTAGAAAATAACCTTTGATTGTGG[G/T]GCA GTTATACTTGGCTAATCAAGAGAAATTCACAAAGTGGCTGACAAAAATTACTGGAAAGCTA TTGGGGAGATCATTCTCGAGAGGTTCCCAACATTGAGAAGAAAAGAAGCAAAAGGATCA GGAGAATAAGCCATCAATCACAGTCGTCCAAGGCCCAAAGCCTGGCAAACCCAC
>Gm07_ 3646081 5	AAGAAATGAGACTGAAGATTATTGAGGAAGCTGAGGAGTATAAGGTGGCTTTCTATGAGAA AAGGAAGCTTAATGTTGAGACTAACAAGGTGCAAAACAGAGAAAGGGAAAAGGTAAAGTTT ACACACAACATATATGTATCTTCTATTTTAATTTAGAAAATAACCTTTGATTGTGG[A/T]CAG TTATACTTGGCTAATCAAGAGAAATTCACAAAGTGGCTGACAAAAATTACTGGAAAGCTAT TGGGGAGATCATTCTCGAGAGGTTCCCAACATTGAGAAGAAAAGAAGCAAAAGGATCA GGAGAATAAGCCATCAATCACAGTCGTCCAAGGCCCAAAGCCTGGCAAACCCACA
>Gm07_ 3751919 1	TTATCGTTTACTGTGCCCTTAATTTTTCTTCTCCCTCTTCAACACCATTCTGTTCTCCAGTTT CTCTATCTAACTATCTACACACAGACAACGACGTCGTCTTTGTCATGTTTAAACTCTTGCG AGATCCATTCAATCCTAAACACTGAAGAAAAATAGATAAATAAATAAAACCATG[T/G]TTGCG GGTAAGGTAAGAGTAGCAATGGGGTTTCAGAAGTCACCGGCGGCGGCGCATACTACTCCT CCGCCGAGAGAAGCAGCCTCCTCCTCCGTCGCCGTCCACCGCGAAATCCTCGTCTCAC AAGTCCTCATTCTCTCGCTCTTTCGGCGCTTACTTTCCGCGCTCTTCCGCGCAGG
>Gm07_ 3751919 3	CATCGTTTACTGTGCCCTTAATTTTTCTTCTCCCTCTTCAACACCATTCTGTTCTCCAGTTTCT CTATCTAACTATCTACACACAGACAACGACGTCGTCTTTGTCATGTTTAAACTCTTGCGAG ATCCATTCAATCCTAAACACTGAAGAAAAATAGATAAATAAATAAAACCATTGGT[C/G]GCGG GTAAGGTAAGAGTAGCAATGGGGTTTCAGAAGTCACCGGCGGCGGCGCATACTACTCCTC CGCCGCAGAAGAAGCAGCCTCCTCCTCCGTCGCCGTCCACCGCGAAATCCTCGTCTCACA AGTCCTCATTCTCTCGCTCTTTCGGCGCTTACTTTCCGCGCTCTTCCGCGCAGGTT
>Gm07_ 3785823 2	CTGTAAAGGTGAGAAAAATTATTCAATGAAATTGGGAGACATCCTTCTCAAATTTAGGAGAAG TAGTAGTCTTAATGGGAGCCCAGATATCTGCACCATTGGTCTCAGCAACTCCAATAGTAAAT GACACAGGAACAAGACATAACCCTCTACTCCTCAGGCTGCATGGCTCCATACCCT[A/G]AA GTAGCAACAATTCAAATTATAACCCAATTCAACACCAGGAATAACACTAGTCACATAAAGAG AAAATAATTCATGCCCAACATCTAAAACCATAAATTGACATACGATAAATTTGTTGACTTTT ACAAAAGGACCTAAAATCATGATTTCTGATTGGTTAGTGATAGAAATTAAC
>Gm07_ 4344233 1	CTTGAGACAACAATATGGAAGTGGCAAGTCAGTCAATGAAGTATTGTTGGTCATTGAGGCT TACAAAGCTTTGCGTGACCGAGCACCTACCCTGCTAATCATGTTGTTGGCCATCTCAGTG GGAGCTTTGCTTTTATAGTTTTTGACAAATCTACTTCTACCCTCTTTGTGGCATCTG[A/G]AA GTAATCATAATCCTACCTACCAAATGTCATGATCCACCACCATATAAGCTTTTCATTTATTTCT CATTTTTTGATGATTGTTCTTGACAGGATCAATATGGTAAGGTACCTCTGTATTGGGGAA TAAGTCTGATGGCTACGTAGCATTGCTGATGATGCAGAATTGCTTAATGGT
>Gm07_ 4430467 2	GTCCAGATCCACCCCCCTGAAGCTGAACCTCTTGAACGTTCTCTTCTTTGGCTGCCCTGCA GCCGCCGCCGCCGCCGCCGCCGCCACATCAACCTCAACGTCTGCCTGCAACATGCACATT CAACAATCATTAAAGAAGCCACCACAAATAGGTCTGAATCTCTTGGGAGAAGAAGAACT[C/G] ACCATTCTCGCCGGCTGCTAGGGTTTTGGCTTCCAACAACAACAACAAGAGAAGGCA GTGGCAGCGAAATAACCTCAGGCTCTTTCTTATGCTGTAGACCTAGATCAAACCTCACACCT

[illegible]

6	TGGTGGCTTGATCCTGAAAACATTGCCATGAAGTCCTCCTTTCCCAACTAGCACAC[T/G]AAGCTCTTCAATGAATATACAAATAATTTCGTTAGACGGTTTGGAAGTTATTTTGGAAAAGAAGGCTTCGGATGCATAAAATAAAATGGTTCCCTTGCTAACCTCTGAGTTTTTCAAACACGACAGCAGTTTCAGCCTTTGCGGGTGTCTTGTGTCTCGGTGAGTACCAACTCTATGC
>Gm08_14760513	CTGATCGAATTATAGTTTCCTCCAAAAACAAAATTGAGTCATCGTCATGTTTTTTAATGAATATGAGTCATCGTCATTTAAGCCTCATGATGAATAGCAGGAAGATTTTGACCAAAACCAAGAGTAGTTGGATGTAGATTATAGATCGAGTTCCTCATCTCATAATGAGCACACGAAA[C/G]GCAGGTGCCTAATAGCTTCAGAAAAGAGGAAATGAGGACCCACACTACCGATGAAGGCACCGTTTTTAACTGTATATCTGTATAAAGTAAATCTGGTCTTTGAAAATGAATGTAATCTATTGTGTATTTTTGTGTTTTAATTTGATATAATTAAGAGAATAAAATTACCTATTCTTAAT
>Gm08_18971181	CTAGAACACCAAACCTGAGGAGAAATTTAAATGTCATAATAAGAAACATGTAACCTACACAAGCACAAATAATGGA AAAATGCACGGTAAACCTACATATATAGGCATGGTTGAACCTACTTGTTAATGACACTTTTCATTGGTTTTCTGATCCAGTCTTTGTGTGCAGTATTTTGCTCC[A/G]CTTCTTTCTTAGTCTGTCTTCCAACGTGGAATATACCAAGCTCATTGGATTACACACAAATTATATCCCTCAGTATCAAAATCCAAAGTTGCTACTCTTTAATTTTCAACAACCTCGAAATGTAACAAACACCACACCCAGAAACAAAATACCCTCCCCCAGGAAGAAGAAGATACAGT
>Gm08_27933899	ACGAGGAGACCACCTTAAATAAAAAATTCATAGTGATTAGATTCTGTAAGTTCTGACTAGAC TCATACTAGTTAAATTCGGTATATTAAGCAATGATTGAAAGATATAGGATTCAAACAAGGCCATCAAAGGCATATAATCTTAAATAACTTGGATAACGATAAGTTTCAACTTTACG[A/G]ACCCAAAAAAAACCTAACATAACATTGAAGATTTCAACACATAAGCCAGTGTTGCATCGGCTTGT TAAATTGGAAAAGGACGAGCATAATTCTAGTCTATCTCTTCTCCCTCACAACACCTACAAG AATTGTAGCAAAGAAAAAGAACACAAAAGTTACAAGATGATAAAAAATAAGATG
>Gm08_37207618	AATAATAACAACAACTTTTCAGTTTCAACGGCTAGTTCCATTTGAAAACAAGATAACACTA AGTACCCTTTTTGTTTTTTATCTCACTTGA AAAAGCGGTTGGTGCACATAACGGTAACCT GCGATTTATTTCTCTTTCCGATTTGCATTTACCATCTTCACTCACCAAATCACT[C/G]TAACA CAACACTACTTTCTTCTGCTCTTCAAGCCAGGTTAGTCAGTTTATGGATTCCCTCTTTTTTAT TTTGAACTAGTTTCCCTTAAATCTTGAAGTATTGACAAGTTTTTCTATGATTTTGATCTTT CTGCTTTGCCCAAACCAATGATTGCATGTAATTTGATGAATCGAAAG
>Gm08_40283689	CCTAATTAACACAAACAAATAAAATAATTTTTTTAGCTTCGGTCCAAAAGAACCGGCTACAG AACTAATTATTGGTCTAGGACTTAATTACACTTAAAGCTTCTCACCTCTTCTAAGAGTGTAT TTTGCGGGGATCGAATCCCCGTGTGTTAACTAACTACACTCATTACATACAAAT[G/C]ATTTT TTTTACTACTTTAGTATTAACAAGTGAACATTTTTATGTACCACGGCATTGCTGGTGGACAC GCCCTTAGCGGAATCGTCCTCCTCGGAGTCGCGCCGCTTTCTGCTGTTACTGGGTCCGAG TCCGTGGCTCGGCCCGGTGGTGAAATCCCCAACTGCCCTCCTGGCCCGTGA
>Gm08_43073940	AGGGAGATTTTGCATATCGATTATTATTTGTTTTTTGTAGAGATATGAACCTTTATTATTACC AATCCATCGCGTCAATTACAAAGGTAAATGAATACAATACAGTTCATAACATCTGTGCATT AATGAATACAGTCTGTATGTATACATAGATAGACAGAGGAGAGAGAGAGA[C/T]JGGA GGGAGGGAGGAAGGGAGAGATTACAACCTCTTGCTATTATTGTTCTTTGCATACTCCCTC GGTCCAAATATTGTGGATCCAATTCTTACATTAGTACTACCCATCTCAATCTGAAAAACAAA AAATATTTTTGAACAGTTGCATGTATACAGTTGATCAAGAGTAAAAGAACATCA
>Gm09_843867	TCTGCTGTTTAATTAATGTTAGTTTCATGTTGAATTGATGCATATATGATGGCTGAGTTGATT TCTTTCTCTTTCACAGTTGTTGGGAGATATGTTACTTGATCGTTCAAATTCAGCCGTAATGA CAAGATATGTGAGCTCAAGAGACAACCTTGAGGATTTTAATGAATCTTCTAAGAG[C/T]ATGG GTCTTTACATAAAACCTTAAATGACATGTCTTATTACTTACCTTCTTGCTTGTGTGTGCT GCCTGTCTTCCAATCCTTATGTATCCTTGTTATATTATCCCGTCTTTATTATTTTGCCGTGA AAGTGCATCAGTTCAGTCAAAATATATAGCTCTTACCCTGCTGCTATTG
>Gm09_3344868	AAAAAATTGCCAAATACGACACCAAGGATCGACAGTGAAGTTTATAATTAACATCACGATC ATGGATGATCAACCACACCACAGTGAACCTGTTTCAGCTAACTAAGTAGCATTTATGAGTAAG AATTGTAGCCTGTGTGTGGCTATAAATACCATCCCTCCCATGCCATAAATCGCA[G/T]CAC AAATAATACTAATCTCCATCTAATAGTGTTAATTTGTCTTTCTCTATCTCTATCGGTCATG GGTGCTTTTGTCTTTGATGAGGAAAACTCCTCCACCGTGGCTCCGGCAACACTCTACAAAG CTCTGACAAAAGATGCTGACACTATCATCCAAAGATTATTGGGGCCATCCAA
>Gm09_6980316	TTTATTGCTCATGCTCTTGTCACTATTGAAGGTCCATATATTTGGTCTGAGTAGAAGTGGGA AATGTCTTATTATCATAATATGCTAAATAGGGTTGACATGCTGTTTACTTGTTAATTTATTTT CTTTCATGAAATGTCAATTTTACATAATTGTCTGCTCTATGGAACAAAATAAAAA[T/C]AGCGA GGATCTGAACCCCCAAAGTATTCAAAGCAAGTGAAGGATGAGGAGGAGGGATTGGACGAA GAAGACGATGACGAGCATGGGGAGACCTTGTGTGGAGCATGTGGGGAGAAGTATGCATCT

	GATGAGTTCTGGATTTGTTGCGACATATGTGAGAAGTGGTTTCATGGCAAGTGTG
>Gm09_25485073	TGAAACGGCAGCCTCCATTGTGTCTGTTAAGGTGCACTCCGACGTCGTTTCGCTCGACGG AAGAGACTTCTTGAAACAGATGCTGAAGTCGGTGATGATGGGAAGCTTCATGTCACCGTT AGAAAATCGAACGCTTCACGGAGGTGTTTTATGATGACGCCGAGGCCGTCTAATCTCA[G/A] JCGGGGCGGAGATTTACAGCCTCAGCTCGTCTCGTAACCCAACGCCACGTGGCTCCAACCT TAACCATGCGGATTTTTTTTCCATGATGGGGTACCAGCCTCGCCACTCGAATTTACCCGCC AATGATTTGTTCTCTTCGCGTGGACCCACTCCGAGACCTTCGAATTTCAAGAAAGCTC
>Gm09_38777811	TTGGGGGTAAAGGATGAGGGTTAGGGTCGTAATAAGGATTAGGATTGGGATAAGGGTTGGG GTTGGGGTTAGGGTTAGGGTTAGGGTTAGGGTTAGGGTTGGAGGGAAGGGGCGAGTAAG AGACGAACTCGTCCTTCCCGATCATCTTCAAGATTGTATGGTACTTTTCAGGAGACAAAC[C /T]AGTAAAACAAAATTATACAAACACCATTATTAATCCTTGCTTTTGCTTAACCTTGACACTCA ACAATTGTGGCATGCTTAATGTCTACAATTATGTTTTGGCATTTTTTAATTGGTGACTACTGT AAAAATATGGTGACTCCTATACTCCATGACCATCGATCCCTCCACTGCTTGCAAT
>Gm09_39791349	AAAGGTGGAGAACACGAAGTTGTGTTCAGACTACTTTGCTCTAATAATGTAGCTGGGAGCG TGATTGGCAAAAGAGGGGCTATAGTCAGAGCTCTGGAAAGTAAACAGGTGCTTCTATTAT CTTTCAGCTCCTTTAAGCGAGCATGCCGAGCGCATTGTCACCATTTCTGCTATCGA[A/G]G TCAGTCATATGAGAATGTTCCCTATGCCACAACCTAACTCAGGCTTATAGAATTTTCAGTGTT TTTTGTGTCCTCTTTCATGTGCATATGTTTCATGCAACAACGCGTAGTCATGCATTTGGTAGA TCTGTTTGAAAACACATTTTGAATTTTATAATTTTATAGTTGATTTAACACTG
>Gm09_44762001	GATTGATATTAAGAAAGGATGCATTTCTTCGGTCCAGTGCCATATACATCATGTTATTACG TAGCATGTCACCATGCTAATAAATACTGAACCGGTTAATTAAGTATTACCTCAATTTTTTCA GCTCAGCTTCAAGAGTTGCCCTTCTTGCTGTTTTCCCAAGCAGCAATAGCTGAGA[G/C]TTGT TTTTGAGCTCTGTTATCAGATCAAATAATTTCTAGAGTTTAGAGATGAACATACATTGGAAAT TAGTAAGGATCAAATAATCTATGAAATGGCATATCATCATCATGGCCTACTAATCACAA AAATCACAAATATCTTAAGACTAAGGGAGGCATGTAACTATTTTTTCTTT
>Gm09_46356061	CGAGGACTCGTTCGTGCAGTCGGCGAAGCCGGCAGAGTTCGCGGCTGAGATCGGCCGGA CGCTGAGACCGGGAGGTTTCGCGGTGTTCCATTTCCGATCCCTAAAGATACGTATAGTTT CAATTCGTTCCCTTGATTTGTTCAATTGTTTCAAAGTAGTTAAGTTGCATGGCTTGGAAG[G/T] CTTTGATTCTTCGATGCCGTATATACGTGAAATTGTTTTGAAGAAAGAGTGTGGTGATGGTG CTGGGAAATTTCGATTTTCGATGATTCAAATGGAAAGTGTTATGTTCTGTTATAAGCATGAT TTGGTTAAGATAGCTGAGCCATTGATTGAGGAGGAGCCATTGAAGCCTTGGATTAC
>Gm10_2447728	GAGACTCTTTAAATTAATTTTCTGTATTTTGTGTTATATGTTACTAGTTCCTTTAAATTAGCCA GATGGAGTTTATGTGTATCTAAATGCAGGGATGCTAATGGAATAAATGGCCACTTGTATTG TTAGCTATCTCTTATGGTAGCAGAATAAGACGTAAACTGGTTCCTTTGCTCCAA[C/T]TCTAAA TTCTTAAATCATGAATACTTACATAAGAAAAGAAAAAAGAAAAAATAAAATAACTATCATT ATAAGTTAAAAGTAGAAGCTTTTTAATATTAGTTTTTAAAAATGGTTTTTACTTTATGCATATT TTAATTTTAGTTTATAAAAAAATATTTTTTTTATTTCTTTGTCATA
>Gm10_4227027	AAAATGCACAAGTACTACATTTGTTTTGAAACCTGCACTGCACATTGCACAACAAGGATTC ATCTTTTTACCTTTTTCACTCGAAAACAACCTCAAACCTTTTTGGGGTCTGATCTGAAGAGCA GAGATTATGGGGTCTTGCTATGGTGAAAGTGACAGATAACATGCAAGTGATAGCTG[G/T]TGA TAAGGACCACATATGGAGGTGTGTTTTGTTCTCTCCCTCTCTCTCTCTCTCTCTCTCTCTC TCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTC ATGGGGCACTTAGTGGCAACAAAAATGGAATGACATTGTTCAATTAATGTGT
>Gm10_4303584	CCCATATCAATGTTATTGCTAGCACCATTGAGTTTTGGTTAGGAGGATTGCCGCAAGAATG AGATTGAATATCACGAGATGAATCTTTCTTTGTAACCTTCTAAGCTATTATTATAAGGAGAATT GCTTCCAACACATCCAGTGGGAGATTTCTTAGTGTTAGAGGCTGCATTGTACCT[T/C]CAGC AAAAGACTTTTAACAATTAACAGTTATCCATCTTAAATTAATTATGAACAATACAAAACAA TAACACTAAATCATCATACCTTGAGAAAGCAGACTGATCAGAACGTCTTAAACATTCCGGT CATCCTGAATTGTAATATCAGCATCTTCAACTCCTCTAAGCCTCTTTAAAC
>Gm10_6625287	TGAGAGTTGAACCATGGGGTCTGAGCGAAGGCACAGGTTTCTGGTTCTAGACGAGGGG GTATTGCGTGAGGCTGGTTTCACAGAGCAGCATCACCCCTGGCTCTTGACAGTGGAGGG ATTACAGGTTGCTTCACTTCCAGCCCATATAAACCTGATGATCCTAGCTCTTGAACCAT[G/A] ACTTTTTTTTTTGTCTGGAACCAAGTTAACTTTGGAATTTTTCTTCTTGAAAATTTCTTA GTAGGTGAAGGCACAGCTTCATAAGTGAATGTTGCATGACACACAAAAACAATAGTTAAAC TAGCTTTGCTTAATGAAAGGTTAGTGTCTATATGCTCCACAACCTCCCTCATTTCA
>Gm10_3337484	ACAATATAAAGGATCAAACATATAAGAGCAACTTGCAAGTTATAACCAAAAATGAAGTGTAT CCATGTTTGCAGTTGTAACCAAGACAATGAGACAAAAAGATTACATGCAGCCACCAATTAG

6	TCAAGGCTTGATTAATACAAATTTTGTTCATACATCTTGAGCTCCTTAGAAGGAG[C/T]ATC AAATAATAAAGACATACCTGTCTCTACATCTAAAGATAAGACAGCATTTTATCCTTTCTTCC TTCAAATACATGAATGATGTCATCACTCCTTGAATGGAGTCATAATTCCTCAACAATCAACT ACACCACTACTGTCATAAAAGTTTAGAATTAGTTTACAGATAAGCATCAGAGCT
>Gm10_ 3992953 2	AAGATTGTCATTGACATGGTTATAAGTAAGATAATCCTGGACATAGTATCCAGTAGTCGCAC TTCCGTCTCCGTAAGTTATGCTGTAGGGGCATGGAATTTAGACTTGCAACCAGGAATTGG ACCGTCATAAGTGGCAGAGCAAACTCTTGGTCACAAGAAATCAACTCTGAAGTTT[A/G]CG AGCCCTTCGGGTCATAAAGAGTCAAATCTATCTGTTGCAACAACGACAACGACAAAAACAC GATGGCGATTTCAGAAAGCTGAATTGAAGCTGTTAAGTGTTACAAAACATGGATACATACACC AAGATCGCTTTTCTGGGACATCTGGAGCACTTAACACAATTCACCCACAGAATGT
>Gm10_ 4148528 3	AACCTCAGTCAAAGTGCTTCAAATGAGCAGTGACATAAGATATATAATTTGGTTGCTCTA ACTTTTGGATGACTTGGACATTTCTTGGTAATAGTAATTGGTGATTTAACAAGTGTTTTTC TCCATTCTTTTAAAACCTTTCTGATATGTGAATTTTGGAGCTTTTCTTTCTT[C/G/C]GCACCG ATCTGAGACTGAACCAACCAAGGTATGCTACTCTCCCAACATAATGAAAGCAAAGTCGAA ACCCATAAAAAAATTCCTCCGGAGGAGTTGAATGTAGAAATCAAATCTGATTTGGAGATAG TTCAAGTCACAGAACCTCCCAAGAGAAAAGCAGGGGTTATTCTTTCTTCTG
>Gm10_ 4264222 1	ACAATGCTTGCATTCTAGTTGATTTGTTTTACCGATGAGCTTTATTTGACTCCTCCTTTCTT CTTCCTGATAAAGTTTCACTTTCTAGCCGACTAATTTAATCTGAATGGGGTGGTGGGCACTA GGTACCTGGCTAAGATACTTGCTCCAGAAGGTTCAAAGAAGTGGCAGTTGGAC[G/T]CTCT ATTGCAATAACAGTAAGTGTGAAAATGCTGTGCAATTTTGTAACTTTCTTCCAATACAATATG AAATTTGTCAGGCTGAGGATTGAGGTCTAACAATTTATGCATCTTCTTTGATATACAACTA AGACATCTGATAATATGATATATATTTATTGGTTGGAATACCAAATTTCT
>Gm10_ 4264222 2	CAATGCTTGCATTCTAGTTGATTTGTTTTACCGATGAGCTTTATTTGACTCCTCCTTTCTTC TTCTGATAAAGTTTCACTTTCTAGCCGACTAATTTAATCTGAATGGGGTGGTGGGCACTAG GTACCTGGCTAAGATACTTGCTCCAGAAGGTTCAAAGAAGTGGCAGTTGGACA[T/A]TCTA TTGCAATAACAGTAAGTGTGAAAATGCTGTGCAATTTTGTAACTTTCTTCCAATACAATATGA AATTTGTCAGGCTGAGGATTGAGGTCTAACAATTTATGCATCTTCTTTGATATACAACTAA GACATCTGATAATATGATATATATTTATTGGTTGGAATACCAAATTTCTA
>Gm10_ 4389694 7	TCTCGTAGTTTCAGGTTACAATATCAAATCACACACACACAAAAAAGACATTTATAGGGTG CAGACTTTTTCTTCAGGCAAATATGACGTGGAACTAATTGCTTTGGAATTTGTGTCCAATC AGCTTTCAAGAGCCTCTCCAGAAAAATCTGTTTGTGAACCATTCTCCTATAATC[T/G]GCAT GACATTTTCCATGCATATGCAATTCAACAATTAACAAAAATCTACATTAGTCTACACTACAAG AGCATTACACATATTGAATAAGGATGCAGATAAGTTAAGAAATAGTACGCAGTGTAATCAT TTAAGTGTAATAAGTTTATGACAGAAATCAATGTTGGCAGTTAAAAAAT
>Gm10_ 4479749 3	AATCTCAAGATTCGTGCATAACAAGGTTTACTGATCGAATACCAAACATGCACATAGGTTTG ATATGATTTTTCTGACTATTTACTTTTGAGATGAAAAGAAAGGAAGGGCACAAATTAAGTGA TTGATATTTCTGTAGAGAATCTGAATATAAGAGAGGATGCATGGATGAAATAAAT[C/G]ACCA GAGAAACAGCATCGCGTGCGGCTAGAACAACGAGGTGAGCGCAAGAGACGACCTTCCAC ACTGCTTGTAACAAGAGATCGAAGATTTTCAATGGTCTGCAAAGCCTCTGGCCTTATCCC TATGTTTGCCGTTGATCCTTTTCATTTGGACTTCCATCCAACAATATCGACCCA
>Gm10_ 4479749 4	ATCTCAAGATTCGTGCATAACAAGGTTTACTGATCGAATACCAAACATGCACATAGGTTTGA TATGATTTTTCTGACTATTTACTTTTGAGATGAAAAGAAAGGAAGGGCACAAATTAAGTGA GATATTCTTGAGAGAATCTGAATATAAGAGAGGATGCATGGATGAAATAAAT[A/G]CCAG AGAAACAGCATCGCGTGCGGCTAGAACAACGAGGTGAGCGCAAGAGACGACCTTCCACA CTGCTTGTAACAAGAGATCGAAGATTTTCAATGGTCTGCAAAGCCTCTGGCCTTATCCCT ATGTTTGCCGTTGATCCTTTTCATTTGGACTTCCATCCAACAATATCGACCCAT
>Gm10_ 5036829 9	GATGAGTTAGGGTATTCAGGTTTTCGACTCACCAAGAGTCTTGTGAAGAGTGCTGCTCAAG CCTTGACGGTGAACTTTCAAACATATGGCGCCCAAGATTCTGCCACTGTGGGAAATGGTGAG GTTCTTTTTTGCATCTATAAGGTATAGTGGGATGAAGAAGAGGGAGCTGCTACCCTT[A/G]C TCACTGGGATGGTTCTTGAAAGGGTAATAAGAAGAAGACAAAGTTTAATGTGTTGGGAGG GGGTTTGAATTTCAAGGCTGGGATAGAGCACTTTTGTGTGCACCCTGGTGGGAGGGCAGT TATTGATGGGGTTGTAAGGGTTTAAGGCTGAATGAATATGACCTTGAGCCTGCAAGG
>Gm11_ 1932902	GGCCACTGATGTTGAGCATGTACAATCAGCTGTCAACTTCGCCTAGAATGAAGGGTATTGT TATTTGTGATAGTTCTGTAGGTAAATTGTTTCTGGGTGATAAGTATTCTCGGTTCAAATATTT TATTGACTCTTTAAGATGTAGCTAGTTACTCGAGTCCAAAATTTATCTTTTTCT[C/T]GTTTT TTTTTTTTTTTTTTTTTGCATTTTGTAGTGAATGGAAATTAGTCTTTGAATAAAAAAAGAAGTTG CAATAGCTTTTATTACAGGCCGTTATTATATGTATGAATCACATGATGTGGTTGGACTCAAT

	TAAAAATCAAATTTTTAGGATTATTTATCATAAAAATCTCTTTTAATAGT
>Gm11_2152220	CCCAACGTGCCTGCTACGTCGGTTCCTCCTATCTATGCTGCTCTATTGACCCCTCAAGGGA GATTCCTCTACGACCTCTTTCTCTATAAGCCTCCCACGTGCCACACCAAGCTCGATCACAC CGGCACTGGCCCTGATTCCAAACCACACGAACCCCTTCATATGTTTGCTGACGTGCA[G/T]G CTTCTGTGTTGGATGAATTGTTGCAAACCCCTCAACAAGTCAGTTTAGCTTCTTCTTCTTTCTA TGGAGTTCCATTTATTTGCTCTGTATCATTGATTCAATTTCTATCAGTTGTTGAGATTTGAGAA ACATACATGGCATCAGCGAGAGAGTTCCATCACTTCCCATTACCAATTAGCTT
>Gm11_2296897	TACAAGAAAGTGGTTGGAAGCCTAGGTGGTTCCGAAGAGAAGGTGAAAATGGAACATTTT GATACGTTGGTGGGTATTGGGAAGCGAGAGAGCATGGAAGATGGGACGGATGCCCTAATA CATTTGGTGAATTTAATGAAGACTCTACCGATCCGTTAGGTGCATCTTAATTTGGTTCA[T/C] GTAACCTTACATATTTTCCAGATAGAGCTGCTTGATTAAAGTTAAAGTTGAACACATACGGA TATTTCAATAAAAAAAACACAAATGAGTTTGCTGTGTCTGAGACACCTTTGTAGGTCTTGA GCTGTAACCTCAGTACGTTTCGTAATGATAAAAAATCGTAAAAAATGACGTACTCAT
>Gm11_8009572	CCATTGCCTCCTGAAAAACCTCCTCTGTTTCTAAACCCCTTGACTTGATCAGAGCCCGCTACA CTCTCCTCAGAAACCCCTTACCACCCACTACTCTCTTGCTCTCTCCCTCGCCATTGGCAC CAAAACCATCAGCCTCATAATCCATAACAAAATCCTCGGATGTCACTCAAAAACCC[G/A]AG ATTCAACAAAGGGGGCGGGGAGGGGAATCAGAAAGCCATATAACCGACATAATGATAAT GATAAGAAAAAACGGAGGCGGAGGAACGAAACATTAAGAATTACTATTCACAAAATAAGAA TTGCAAGTTTCAACAGCAATAAAGAATAAATTAATTAATGGATGGCCGATAGGAATTA
>Gm11_11628041	CACCGTTTGATTCCTCTTCGCGATCATGGATCTGATCTGGATCACGAAATCCGGTATTGCA AACCTGCGTTTTTTGTTTCTAATTTACGCTTCTTTATTTTTTTGATTTTGAAAATTTTCTTAGGG CTTGATTAGGCTTCCTGTTTCTCTAGCTGTGAAATGATTGGATCTTCTTCTTGC[C/T]CTTGA AGGAAGTCGCATTTGGTCTTTCTTGAAAATTGATCTCTCAATTCATTCAATTCTATTATTACA CTTTTTTTTTGCTTAAATTTCTATACAAACAACGGCTGCATTTACGCAATTTCTATTATCAC ATTGCGTTTCGTTTCTGTTAATCACGTTTTATTAGTTTATGAGCTGCCGC
>Gm11_11628062	GATCATGGATCTGATCTGGATCACGAAATCCGGTATTCGAAACCTGCGTTTTTGTCTAAT TTACGCTTCTTTATTTTTTTGATTTTGAAAATTTCTTAGGGCTTGATTAGGCTTCTGTTTCT CTAGCTGTGAAATGATTGGATCTTCTTCTTGTCTTGAAGGAAGTCGCATTTG[A/G]TCTTTC CTTGAAAATTGATCTCTCAATTCATTCTATTATTACACTTTTTTTTTGCTTAAATTTCT ATACAAACAACGGCTGCATTTACGCAATTTCTATTATCACATTGCGTTTCGTTTCTGTTAATCA CGTTTTATTAGTTTATGAGCTGCCGCATTTATTATAAAGTAATGATG
>Gm11_17315140	CATCAAAATGATCATACTCTCATATTTTTAAAAAAAACCTACAACCTTTTCACTATTATTGTC TTATGTCATATTACGTTCCATATCATTATACATGCAATATAACTAAATAAAAAAAGAAATC CTATTATCAAGCATATAAACTGTTCCATTATGAGGATGATAAAACAAACCATAC[G/C]TACCAAG TTAGGTAACATTGCACCAATAGTATCGGGGTTTACAGGAATTGGTGGCTCATACATGTTTT GACTTCGGAATGAATGAAAAAGTGGAGATGAAATAGCTGATTTTTTGGGATCCAACAAAGA AATACACACTGAAAGAGAGTTGACAAGGCTAGACATTGATTGTGAATCTT
>Gm11_18487772	GTTGAAAGTTGCCACAAGTTGAGTTTTGCTTCTGTAGTAAGGATCCTGATGACTTCTTCAC TGTTGCCTTTCTTCTCCTTGACAGATTATGAAAGAACATTAGCTTCAGTTTGTGAAAATTTTA CATTGATCTCATCACCTCATACCTATATGAGGTCACCAATCCAACCAAAAGCT[G/T]CTCAA GATCAAAACACAATCACCATTAGCACTCCTCAAAGTTCTCATCAATAAACACCATTCAGTTGC TACAAAACACTAATTCTTAATAATACCTATATCTATATGTATTATTAATCAAAACATTAAAT AACTGGTGATTTTAACACAAATTTTATCATTATTTCAATGCACCCTTTA
>Gm11_18487773	TTGAAAGTTGCCACAAGTTGAGTTTTGCTTCTGTAGTAAGGATCCTGATGACTTCTTCACT GTTGCCTTTCTTCTCCTTGACAGATTATGAAGAACATTAGCTTCAGTTTGTGAAAATTTTAC ATTGATCTCATCACCTCATACCTATATGAGGTCACCAATCCAACCAAAAGCTG[T/A]TCAAG ATCAAAACACAATCACCATTAGCACTCCTCAAAGTTCTCATCAATAAACACCATTCAGTTGCT ACAAAAACACTAATTCTTAATAATACCTATATCTATATGTATTATTAATCAAAACATTAAATA ACTGGTGATTTTAACACAAATTTTATCATTATTTCAATGCACCCTTTA
>Gm11_24415494	CAAGAGCAGGGCCACTGAAATAAAGCTGAGTCGGAGAAATAGCAAGGTCAAGGGTGCTG GAGAAGGGCGGACGCAGAAAGTACACCACCGCCACCAAGTGACGGTCATCACTGGGTGGTC ACCAGAAGAACACACTGCAGGATGTGCGACCTGTGGCCACACCTCCTATTTGTTAGGG[G/ T/G]AGGCGGCGTGAACCTGACTTGATCCGGGTATTCAATCACCCCCACAAAGGTTAAGGA GTCAACTGAACATCTCAGCCATTGGCGGGAATTTACCCGCATCCGACCCCCAATTCTTGT TCACCCATGGAGTAGGTTGCTGCTTGACGCTTGTGCTTATTGTTTGAGAAAAGATGACTTG
>Gm11_2655975	TCACCTTGTTTTATAATGCTAACTATTTCTATGTCTGACTAGGGTAAGAACAACCTAAATTA GCTCAGAAAGCAGACACTAAGGCCAGCCATGACATATTTGTAGCTAGGTTGGTGGATTTC

8	CCTTTTCACATTCTTGTGTAATACTATTGCCTTGATTAAGCATTGTTAGTGGCATAT[C/G]CAGA AACAAAGGTATAGAAAATGTGATAACTATTGACTTACATGGGCAGCACGTAAAAACAAGCAAT GAGAATGCTGAAACTTCACCTTCTGTTTGGATCATATGTTCCCTGTAAGCATTITTTGAGGAT CTAGGTTCTCACATTTTCTTGTGACACTAAAACCTGCATTGCATTTTTCATTTA
>Gm11_30156178	CTTGCGAGAGGATGAAAAATGAGACGAAACATTTGGAGGAAATGGAAGGAGCAAAGAGTCAT CTCCATTTTTTTGAAATGGATCTTCTTGACATCGACTCCATTGCCGCTGCCATAAAGGGTTG TTCCGGCGTAATCCACCTTGCATGTCCTAACATCATTGGTCAAGTCGAAGATCCC[A/G]G TTAAAAATATTTAAAAATCACTTAACACACTCCAATATTTACATAAATGGCTTAATTACTAATGT ACTTCATAAATCCACAGAATCTTTATCTCTTAAACATTACTCTTTTAACTCTTTACCCACACAC TCITTTTCTCTTCTCATCCATCAAAAGTTATTGATACAATTAATTTGATATT
>Gm11_35282159	TAATCACTAGGGTTTATTATCATCTCCATCCATGGAGACTAACCGTTTTCTCTCTTCTTCTTCT TTCACCTTCCACCTTCCCTCCTGCTTCTCAGGTACCTCGTTGATTTCATTCTCCGATTTGCG ATTTCTTCACAATTTTAACTGTCAATTTCTCACTGTTGCGGTTTCTTTTCGAG[A/G]ACAATC GAGCTCCATGGAATCAGTGCCTGATCTTCAGAACACTATGTACGCTTCCATCGACGGTTAT CCTTGCGTGCGCCTGATGAATCTCTCTGGCACCATCGGTTGTTCAAGTGAGTCTCAACCT CTCGCATTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT
>Gm11_37298431	CGTGCTCGTCTCGGCGGGGAGCCCCGACACCACCCGCACCTCGCGGACGGCGCCGACG CGAATGCCGTCGCCGGTGATGACGTGGCAGCTCCTCACGAAGTTCTTGATCCCTGCGGG TTGTCGAAGCGCCGCACCACCGCCCAAACGGCGGAGACGGGGGCGTTGATGTGCTGCGT CA[C/G]GATGGAGCAGCACTGGTTGGGGTCCACCACGTGCGCGTGGTGGAGCGCCACCGT GTCGGGGGCGCGATTCTCTGCCGGAAGAGAGAGAGGGAACCACGAGGCGACGCGCC GCCGTGGAAGAGGGCGGCGCTTGAGGCTGCTTCGGACAGTTTACGCCGTTGGCGATGGC GACAGTAGGT
>Gm11_37298433	TGCTCGTCTCGGCGGGGAGCCCCGACACCACCCGCACCTCGCGGACGGCGCCGACGCG AATGCCGTCGCCGGTGATGACGTGGCAGCTCCTCACGAAGTTCTTGATCCCTGCGGGTT GTCGAAGCGCCGCACCACCGCCCAAACGGCGGAGACGGGGGCGTTGATGTGCTGCGTCA CG[G/C]TGGAGCAGCACTGGTTGGGGTCCACCACGTGCGCGTGGTGGAGCGCCACCGTGT CGGGGGCCGCGATTCTCTGCCGGAAGAGAGAGAGGGAACCACGAGGCGACGCGCCGC CGTGGAAGAGGGCGGCGCTTGAGGCTGCTTCGGACAGTTTACGCCGTTGGCGATGGCGA CAGTGGTGG
>Gm12_4386073	AAAAGCATGCTTCCAGCTCTATGACAATAAGAAAAGAAGCCCAAATAATGCAGTCAGAAAAA ATTATAAACCTGAACAACGTAGTTGCCATACTGACTGACTAACATGCTAACTGCTCCTA AAATTTTCACTCAATAACTTTCTGTTGTGTTGATAGGCTCTTTGCAGTGTTCCAGCACT[C/T]GCT GTGCGGTAGAACGTATTTTATATGACAAATGATCTAAGGTACATTATAAGAGTAACATAAGC AGTGAAGAGAACTTACCTGTATCACCCGGCAACCATATGGATGGGTTGAGAGTGTACAAC TTGATCAAAAAATGTTGACACAATAAAGTGGATTGCATCTTCAGGGACACATTC
>Gm12_16133337	TGTTGCTGCAGCTTCTCCTGCCCTCAACAGCCTGCTATTAGTAGCACTCCACCACCACCC CGACCACCAGCTGGTGCCAAACCTGGCCACCCCTACCCTACTGCCAGCACTAGCTGGT GCCAGACCTGGTCTCCTCCACCACCTCCAGCCTCAGTTGGTGCCAAACCTGGTCTC[G/T] JACCACCACCCCCAGGACCAGCTGGTGCCAAACCTGGTCCACCTCCTCCACCACCCCCAG CTGGTGCCAAACCTGGTTCTCGTCCACCACCACCTCCTCCTATGAGTGGTGTGGCTCCTCC TCGACCTCCGCCCCCGTTTGGTTCAAAGGTGCAACGTCCCTTAGCTAGTGGATCAAAAGC
>Gm12_17533119	GTCAAAGGGCACCACCAAGTGTGTACAGCATTCTTCATTGCGTTCTGAAGCTCCACCTCGTCG GCTTTGTCGAGTCTATGCTCAACACCACGTTCTTTCTGCACACAAGTTGAATGGGGTG TCTTGTCTTGAATCCAGCCACCTCAGCACATCAACCACTGTGAACGATAAATC[T/C]T GAGCACACAAAAAGCAAAAAATAAAAAATTGCCAATTGTCAATCACCAAACCCATGTCCAAT TTTGTGGTATCTTAAGATAAAAAAAAAAAAAAAAAACATGGTTTTAAAAAATAATCAGAGTTTTAG CAGTAAAGTCAAGGATTTTGGAGTATCTGCTACCGTAATTGTGATCACATGGTA
>Gm12_18823992	TCAGCGACACTGTGATCTGATTCTCTCCCGGCTGCATACCAACAAAAAAGGATCTACAAC ATATAAAATGAATCTTTAACATTTTTCATCATCCCTCAAGAATAAAAAATGGTAATATTCTAA ATGAAAGGAATAAAAAACTTAGCATCAATTGAACAATTGAAAGGAAGAAAAAAT[A/G]CTTA TCCATATCAATGGCAATGCCAGAGTTGAAGTGTGCTCCATAAGATGCCACATCCCTGGTT TGGGTTTTCGAAAAGGATCATCTTCTTTTGTGTTACTTTACCTGATTTACTACCTACCCAC AAGCTATAAAAACTGTATAGATAAAAAATAGATACTAGGTCAGCTCTTCTGT
>Gm12_24557321	GAATGTGAACCTTGAAGAAGACTGTCAACTGAATAAAGCCCATCCAAGCTCTTTAACGTGT TGCTGATCTTCTATTATCTCCAATCCATTCTAGTGGCATTGGGAAGCTGCCCACAGTATCT GGTCCAACACTCACAGTCAAAGTGCTAGAGTCTGCTGAGGAATTGGCAACAAAGTC[C/T]TT

	GTAGTACGGCACAGCCAAGTCATTAGTTATGGATGAGATGTCAAAGCTTTGAAGAGCTATG TCAGAATTTACGAACACATTGAACACTAAAGTGTTGAGTGACTTGCTCATAATATCACAAAA GTGTGCCCGGATGAAATAGGAGAAATTTGGATCCACATTGAAGACCCAAGTAATG
>Gm12_ 3800191 1	ACACAACACAAACTATCTTAACAGGAATTATATTATGTACTTACCCTTTTTGGTGGTGAATAA CCTGCTTCTGCCAAAGTGATTAACACGGTTCAAAGAGAGAAAACAAAATAAGACCCAATTTA GATTACGTGAACCCCATTAAGTAAATCCCATCAAGTTTAGCTTTTTGGTGTGGCC[A/G]CAA TGCAAGCAAACCTCTTTCCAGGCTGTTGATTAGGCTGTCAATTACAGTGGGAAAAAAGTG GACCCACAAAGAGAAAAATTAATAATTAATAAAGTGTTAACAAAACCTCAAACCTGAAAGTTG AAACTCCCATAGAACCAGCTGTTACTTTAAGATTCTGAGTTTTCTAAATTCGGA
>Gm12_ 3858885 2	AGTGGAGCATTTTTAATTAGATCAAGGAAAAATCTATGCATTTTAAATTATGTCATACCTGGA AAGTTTGATACAAAACCACTCTATTTTTATATCCTCATGTTCTGTGGTTGGCTCATGATCC ACTGATAGTTGTTTGGCCACACCCTTCTTACATAAGACAGCCCGAGAATCCCC[G/C]TGTG GCTACTATTAACCTTATGACATTGACTAATATTGCTGTAAGTGCAGTTGAACCCCTCTACCTA ATTACCCGACATCTCTAAAATACTAGAATCAGTTTTACTGTATGCTCTCTTGACAGCATCT GCAGGTTCTTTCCAGAAGTCAGGCTACACGAAACAAAAACATGATTGAACT
>Gm12_ 3949380 8	ATCTGGTACACATTCCCAGGGTGGATCTTCGGATAATTGGTCTAGGCCTGGCTTTTTCTGAG CTGACAAGCACTTCCCTAAAAGAAAGATGGTCATTTGACAGTGAGTCGTTTGGTTTTAATTG TGAAAGGCCCGTTAGATCCAGTAGTCGGTTTTCAAATCTCCAGTTGATTTGCAAA[G/T]AT GTGGTGTGTTGCTCAAACTTTTAACTGAGAAATCTTCTTGGGGAACCCAAAAGATAATTGCA AGTAATGACCTTTCTGTAGTTTCTGTCTTATCTGTGGACACGTTTATCATGCTGAATGCTT GGAGAGTTTGACATCTGAAGTCAACAAGTATGATCCATCTTGCCAGTTTGCAC
>Gm13_ 6166640	TAAGGCCTTTGGAATCCGGGAAGCAAATATTAAGTTCCATGGCATCTTCCACAAAATGTT CAGGCATGGTAGAAAATTCATTGGGCAAGTTGGTGGTAGAGGCATCTTAAATCCACCCAC CAAGCCAACCAACCAACAATCATCAAGCGGTAGAAGGAAAGCGCATTTTGAATAA[C/T]TG TGTTATCCTAAGGAAACAATGTGAAGCCAGTAATTCAAATGAGAATATTAATAATTTGAAA CTTTATAAGCTAGAAAGAACGCTTAAAAGTAACAAGAACATGTCTAAATCAAATGATTGATG AATCAAGTATCAAATTTAGTGAACATTTAATTTAGAAGAACATAATCTTTGTA
>Gm13_ 6388118	ACACGGACATGGCACAGAAAAGCTGAACATTACCTTTCAGTACATATATGGAACATAATAAA GCAATAAAAGAACATTACAGGATCATAATTACTCTAAAGTTAGTCAATATATAAAAGTGATAA CCTGACTCTTCAGATATGGCTGCAATGTAACAACAAATTGGGAAGGGTTTGAAG[G/T]GGT GCACAGAGAGTTGGGTCCACAAGACAAAATGCATGCAAGACTAACACATATGGAAGTGCA CCCACTTCCACTCCATCATTGTTTCTTTCGTCAACCTGATTATTAACAAAAACTGTCATTTAT ATATCAATGTAGCCAAGCATAGAAGTTAGTGGCATAAACAAAATACTACCATA
>Gm13_ 6635251	TTATTAAGTATTTAATAAAGATAATAGAATAATTTGATATTTAATTAACACAACCTTAAAGAGA AGGGACTGAAAGTAATATAAAAAAGAAAGGATATTTGATGTAATTTAAGAAAAAACGTAAA GAGTGTGAATGTAATTTATTCTTATAAATATCAGTTTTTCTTATAGCTGTTAC[T/C]TAGTGGG TTAAGCCTCGCTTCTAACGTCACCGACTCAGTTGCTGTGTCAGCGAGTGAGTCGTTTCT CTGCAGCAACAAGGTACGAAAAGAAAAAATACTTCAAGATCTCACTCCTTTCTTTTTTT CCATTTGATCCACACCATGTGCTACCTTATTTTCTTTCTGGGTTTTAA
>Gm13_ 1101331 2	TTCTAATTTGATGAAGTCACCTAAATTTGGAAGAAGTGTTTCAGTTTTCAAACCTAGAATCT GTCAATCTATAAGATCTGCTACAAGCAAGAACTAGCAGGCACAAAATATTGCTTGGTCCAT GAATCAATTCGTGAAATTGATGTTAACGAGCTAGAGTGAAGCTAGATCTGAGAAT[G/A]ACC TAACAAATATGGAACCTTTTACCTCAAATCTACCATTCTGTGAAGCAAAACCAACAATG AATTAGAAAATTAATCACAAGCTGAAAGGAATATACTTTAGGGGGGAGAAATGATTTAC ACCATCTGCTGCACTTGGAAGGAAAATGACACAGAGTCATGAAATAAAAAA
>Gm13_ 1558375 2	CGATGTTAATAGTAAGACTAAATTTCTGTGTATAGACAGTCGGATCTTTGTCATCTATTTA AAGTTTTGACTAGGGGCTACTGTATTAGTGGTCTTGTATTTTGTGCGGGGTATGCCCC GTTTCATTGTTGATTATTTTAGTTCAATATAATTTACATTTTTCTCAAAAAA[C/T]TAAAAAG AAAGAGCTCTCAGTCTGTCAATCCTTACTATGTCTAGAACTGGTAAGTTTCCCCATGTTGAG TCAAATTAAGCCGCAGGCTCCACTCTTGGTGGTGCCCTCCGTCAATTCCTTTATGTTTCAG TCTTGCGACCACTACCCCTGGAACCAAGACTTTGATTTCTCATAA
>Gm13_ 2386133 6	TGCAAGAGACAATAACATGACAAATAGTGCAATGGAGAAAGAGACCGACACAAGCATAAT TGGCTCTGATTTCTGTATCCTCCTTTCCCCCAAAATCCTTGGGAGAAAATAGAGAAAGAGA AACTTTACCAAAAAGAAAAAAGGGGTGAGGAAAAGAGTGATACCCTATGACATGC[A/G]G ACCACCATCATGAACAGCTTGGGAAACCAAAACCCATTAAACCAATGCTTCTCTCCATAG ACCAGATCAATGTTCTTGAGACCTTCAGATACCATTAAATAACAACATTAAACAACCTAGAA AGACTAGAGTAACAGAAGCAAATAAGAGAGTACATGATAAAATGATGCCAAAAACA



>Gm13_25653434	TACTGATGTTCAAGCCATGAGAATGTCACCTGTCTATCCTGAGAACCGTTTGCAAATTGGTC AAGAGCTTACACGAGGTCTTGGTGCTGGTGGTAACCCTGAGACTGGAATGAATGCTGCGA AAGAAAGCAAAGAATCAATAGAAGAGGCAGTCTATGGAGCCGATATGGTCTTTGTTA[G/C]T GTAGGTTTCATGTACTTTTATCCAATTATATCTTCTGCAAATTTAATCATGATTTTGCAATTATG GATTATCTTTTTAGTACTGGTCTAAATGTTACACCATTAGACATGATTTACATGCTCCATTAT AATATCGAAATGTGTCAATCAAGGACCAACTCTCGGAAATGCTTAACCTGTTA
>Gm13_29099729	CACCAGCAGGCTTGAACCTTTGGAGTTCTTTGAAATTCAGCCATGGTTTCACCCACACCTT GGCTTCATAGAGCTTCTTCTCACCAGCCTCAATAGCCTCAAGAGTAAGGTGATGCAGGGTA CCAGCAACAACCTGTTCTGTGCCCTAACACCCTAGCAAACCTCCAGAAGCGAATTC[G/T]T CAACACCAACCAACACAAGTCAACACGTTTAGTTTACAATTACATAAAACATTGTTTCATCAT ATTGAACATAAAGAACTACTGCAATCACAATAACAAGAAAGTTCAGCAATCCTAATAACATG TCAGTGCATTAGTGGGAGCATAAAACCCTCACTAGCAACATATAAACAGATGCAT
>Gm13_29100660	GTGTATGGCCATAACATGACAATTCACGAATGATCAAACACTAGCATAAGCAACCTGGTCT CGTGCATTCAAAACACGTAGCAAGTGGCAATGAAAAAAGATTAAAAATAAAACAATTCTTT TATTATTTTTGTGAAAAATAAAATAAAGGAGAGAGAGAGAGTGTGTTTAGCAGCA[T/A]CCT GCTTCTTGTTGTGTTCTACGGCGAATCGAGCGAGGGCTTCGGTTTCGACGCTGTTTTG AGAGCCTTGGGAGTCGTGTAAGCCTCCGATCGTGGCCATCGGCGAGTGGTGGTGGTGGT ATTCGGAGCAGTCCCCGGACGAAGATCGAAGAGCGAAGAAAATGAAGAGAAAGAAGA
>Gm13_30454510	CCTCAGTTTTCATCGATGGAGAGTGAAGCCTTAACCTCTGCTTATTCGGTGTCCAGATCTCG TTGTTGCTTCCGATTCTTTGATTCTTCGCTTCTTTCTTCTCACTTCACATTCCGATTCTCG CGCAGATTCTTTTGGTTTCTAAACATCATGATCGTCACTGCCTCCATCTGTAATC[T/G]GTGA TTTTCTCTCTTTCTCTCAGCTCATGAATTTTCTTTTCTTTATTAGCTGATGAATTTACAT GATCATCGAGTAAATCACTTTTTTCTTCTATTTTTATAGTTTGAATGTTTTTCTATTTTC AGTTTCTTTTTACACTTGTTTTGTTATTATGTTTAAATCTGTAAAGAT
>Gm13_35154543	CTCTATGCTATTAATCATGTTATGATATATTGAAAAATTGGATTCTTACAACACAAGTTTG ATGTGGATATTTACTATGTTGTGTCCCATGATGAGAAAAAGCCAGCCAATTCATATCCATTT TTCAGTTAACTAGTAGTTTCTTTATATAGTTTGTGATGAAGCCTTCATTTCCAA[C/G]CAGGG GGTCAGAGTTGCCTCAAAGGCAGTCTCCACGAGGTGCACATCAACATCGGACATCAAGTT CTGACTCTGATCCATTGCATCATCGGCCGATTGCTGACCGAAGTCCAAAGCTAGGAGATCG GCGTTCCCCCAGAGGTACTCAATCTGAGGGATTAATCAGAAGAAGCTAGGAAC
>Gm13_37094440	CACTGCTGTAAAGGTAGTTTCGGTTATCTTGATCCTGAATACTTCAGAAGGCAACAGCTTA CAGAGAAATCAGATGTTTATTCATTTGGAGTAGTTTTGATGGAAATTTTATGCACAAGACCA GCCTTGAACCCTGTCCTTCCAAGGGAGCAGGTTAACATAGCCGAGTGGGCAATGA[A/G]TT GGCAGAAGAAAGGAATGCTTGATCAAATTATGGATCAGAATCTGGTGGGGAAAGTCAATCC TGCTTCTCTTAAGAAGTTTGGGGAGACAGCTGAGAAGTGCTTGGCCGAGTATGGAGTGGA TCGGCCGCTATGGGCGACGTCTTGTTGGAATCTTGAATATGCTTTGCAGCTTCAGGA
>Gm13_38425493	CCTCTTCATCCTCTCCATAAAATCTTCCACTCATCATATATAGTCTGCAAAATAGTACAG GGTTGATATCCCATCTTCATTCCCAACTCTGAGTTGTTCTTTACTGTAAACTACTTCTTCACT GTAATAAGGGGTTAGAACACTGAAAGCCATCATTTTCTCAACTTGGGGGGCAT[A/G]GGGC ATGTTTCATAAAAGTGAATTGGTGAAGAAAGAAATCCTTCTTCTGGCTTCTAGATTAACCTGG GATGTTTTGCATTGAATCTCTAGATGTAAGAATAGTATGCAAGCGCCGAATCTGTCTGTAGA AGTTCTCATTGATAGCCTCGGGCAACTGAGTCGCATTCTCGAAAAGTAGAAC
>Gm13_38806205	TTGGCATAATGTAGCAAGACAAATTTGTTGAGGAAAGAGAAGAGGTCTAATAATTGAAGGA AAGAAGCAATGAGAAATGTCGCCAAAGCAGTCTAGGGAAGCTATTGCTTTCAATATTACAT GTAAAGCATCTGCAACTTGTTGAAGATCTACCGGTACACTAACTAAACAGCGGTATA[C/G]T TACCTTCAATCCTTGTTAAGGGTTCGGTCACCAGGTCCATGTGCCTTGAGCCAATTTCTCG GCAATTACATCATTTTCCAACGGCATCCTTGGCAACAGTACCGCCAGCTGAAGAACCTGAAC TGTAATTTTTTTTCCCAGTCAGCTTAATATAGATGAAGAAACTGGGATTGAATAATA
>Gm14_314767	TGCATTGCCTTTTTTTTTTCTCTTCTGCTTCAAACTATTGGATCTGATTTCAATATGTTGAT ATGATTATTATTATTATAATTATGTGCTTGCAATTTGCTGTTTTTTTTCTTTGAAAATCTCTG GCCTCTCATGTTGAAATGTTTATTCCATTGTAGATTTCTGTGTTGAATGCAT[C/T]ATCAGAGT TAGGTTTTATTTTTGACATTTGGTGTAAAATGGGGGTGCCAACTTGAGAGAAAACCAAGTGATC TACTGGGTTTGCCTTTTCTGTCATTATCAATGCGTTTGTACAAATTAGCAATGCATTTTATCTT CCGGGAAGCTACATGCATACTTACTCAAATGGAGATCCGATATATGC
>Gm14_314770	ATTGCCTTTTTTTTTTCTCTTCTGCTTCAAACTATTGGATCTGATTTCAATATGTTGATATG ATTATTATTATTATAATTATGTGCTTGCAATTTGCTGTTTTTTTTCTTTGAAAATCTCTGGCC TCTCATGTTGAAATGTTTATTCCATTGTAGATTTCTGTGTTGAATGCATCAT[G/T]JAGAGTTAG

	GTTTATTTTTTGACATTTGGTGTAAATGGGGGTGCCAACTTGGAGAAAACCAAGTGATCTAC TGGGTTTGCCTTTTCGTCATTATCAATGCGTTTGTACAAATTAGCAATGCATTTATCTTCCG GGAAGCTACATGCATACTTACTCAAATGGAGATCCGATATATGCCAA
>Gm14_ 357009	TCCTTACCAAAAAGTGAAACACGTATTATAGTGAATATTAAGAAGATTAAAAAGTGTGGG GGCAATTAGTAAAATTGATCTCTTAGAGAGAAGTGTGGAAATAATGGAGGATGGGGTCCAT CTGGAAGCCAATCGAAAGCTGGCATTGATTCATAAACCCACAAGACACAAGACAGA[T/C]CA CAGAGCACACTTGTCCATTCTCCAAAACAAAATGAAGAGCCTCCTCTTCTCTCTCTTCTCT TCCTCTTCTTAGCCTCCTCCGCCGTCGCCACCGTCCCCGCCACAACATCACCCGCATGCT GGCCGCACACCCCGGCTTCTCCACCTTCAACCACTACCTCTCCGTCACCCACCTG
>Gm14_ 2810880	TATGTATCTTGAGAGTTCAAGTGGACATTCAAATTATGTGCTAGTATCAGTATTTTAGTGAAT ATATGCTGCTGCAGTCCGGCTTGTGTTTTCTCTTGAACTTTCTTGTGTAATTAATTGATGA TCATGACAATCTAGTGCTGTTTTAATGTGAGATAAAAGATCATTTGCCTTGGG[A/G]CTTTGT GTGGCCGGTTTTATTTATTTTATCTAAGAGGTGGGGTTTTGCTTTGTCATTCCGGAATTGAG AATTGACTATGGCTTCTTTTGAAGGTTGTGTGCATGTGTGGTTTCTATTAGCCTGCTGGTA TAAATCAAACAGTTCTATTGAAAAGTTTGAAGTTTGCTCGTACCAAAACA
>Gm14_ 4182470	TTGGGGGGGAAGACAAAGAGAACCACCCCAACATGTGAAACAAAAATGCCTTCTTTCTGGC CCACCCTAAGGAAATTAACCACTCTTTGAAAAACAACATCAACGGGAAATACTCGTCATCTT CTTGGGAACAACCTTCTCCTTCTACCTCTCCCTTTTCTCTCTATCTCTTCTCTCTC[T/C]CTCT CTCTCTCTTATTAAACAAATTGATACCCCTTTAATTAGTATTCTTAATTACTCTTTCCATTT CCATATCCTAGCTAGCTACAAATATATATTATGGTTTTAGTTCTTTAAGGCATCAATCTGCAG CATAGTCATGAACAAGAAGAAGAAGAAGAGGTTCTGTGGATGGCACT
>Gm14_ 4345820	CAAGGTATCGCAACATTGAACTGTAAATTAATGCAGAAAAATATCTCCATATCTGACATC AAAAAAGGAAACTCACTCAAACAATAATTAATCTACAGAGCCAAATGCTGAACCGAAAAA AAAAATCACAAATCAATTAGAATAACTTTGCGAAGAAGCAAAGGGAGCGAGAATT[A/G]CTA GGTTTGAACATATAGTTACTCCTGAATCCTGCCAACACTAGCAACCTTGCCCCAAAATTCCTG AACAGAAATATGACAAACAGTATAAGTTGAACATATCAATTTGTAAACATGTAAGAACAATCA CAATCGATGAGAAAAATCAGAAAAATAACAATCAGAAATTGCAACGAGATTTCTA
>Gm14_ 5037341	ACCGCTTTCTCACACCTAGAGGATGGGGCTTATCAGATTTCTCAGCTGGACAGACAAAATC TTTCAAAGGCCAGCTTATAGGTCTCATTTATGCAACTCAGTACCTAAGATTTCTCTTTTACT TCTCAACTCTGTCTTCATTATCGTGAAGTTAGTTTCTGGATGATGCCTGACAAGG[G/C]GAG AAATCAATACATATGTATGCTTTAGACACGTATCTAGTGCATTGCCATTGTTATTATTTTCTAG ATGCTAAATTTTACTTTAATCATAACAATATCTGTTGTTCAAGTCACCTTGAGCTTTTCTTAT GTTTGACTAATAAACAACACACAAACATGAATTATATATGTTGACATTT
>Gm14_ 4735439 6	ATTCCTTTGTTCTCGTCCTTTTCACTAACCATCCACTGTTGTTTGGATTCTTTCTTTATATATA GTTCTTCTCTTCATAGTGAGTGAGAGAGAGAGAGAATGGTTTCCAGGGTTCACAAGAGA ACAGCACTGTATAGGAGCATACAACAGCTTCGTTCAATCACTAACTCCCATGCG[G/T]TGAT TAATTACTCCGAAACATCACTTTTTTCTCCCTTTAATTATCTGATATATATACTTCTTCTCTTT TAGTCAAAGCTTCTCCACTTAATATTATATTTGAATAGAATTACCACTAACTTGATATATA TCAATATCCTTAATTTATTAATTATTCTACTTATATTCCATGAGCTAAA
>Gm15_ 28475	CTAAATCTGAGCCTACTGAAGTTAAAGAAGAAGTTCTGGTGCATTCAATTCATGAAAATCTT AGTGAGACAAAAATGGATGAGGATAGTGCACATGATAAAATGCCCACTGGTAAGCCTGTCA CACATACTGAGCCTGCTAATATAGCTAGGCCAGAAAATATTAAACTGAGAAACAA[C/T]AT GGACAAGATAAGCAAGAAAATGTGGACCAGCCATCTGATCATGGGGCTGGAACCAAGTCT GGTAAGCCAAAAATAAAGGGGGTCTCATTGACTGAGTTGTTTACTCCTGAGCAAGTCAGGG AACATATCACTGGCCTCAGGCGATGGGTTGGCCAAGTAAGACTATTTATTCTTGTTG
>Gm15_ 222394	ATCCTGGAACCGCTGCAAGCACATGCAACACCAAAATAAACCACTTTATGATTCTGTTTATA AGTAAAATTTACATGCATGCAACATGAGACATACCTGGTACTGGGGGAAGTGATCCGGCT CCGGCTCCGGCTCCGACTCCCGCATGTCTTCTAGTGAGTAGCAGGAGGAACAAGAG[G/T]A TAGAAAAAGCAACTACGGCTACAGTAACTGAAAGGGTGAGGGACTCCATAGCATCTAACTT CTGATGGAATGAAGATGATGAAGGTGTGGTGTGGTCAATCTTGGAATTTGGAAGCTATAGT TATTAGTGAAAGCATCTTTCTCTTCCAGACTCCAGTCGTGTCGTGCCACCAATCAC
>Gm15_ 440713	GTTGTTATTACTCGAGCAGGACAAGCCACACACAAAGTTATAAAGTGCAAGTGTAATATACT CATAAAAGCATAAAGATGACCACAACAAAGTTGTCAAACCTTCATAAATCAAAGTTCATAC TTCCAACACGATTGATCCACAAAAAGAGAAAAAAGGATGAGCCAATGCACGAC[T/C]ATC TGTTTGAATAATTTGATACACTAGTACGGATTGGTATTGACGTTTCGTGTGATTCAACAACAC TTGCCTTAGTTTTTAACTAGATCATTACCTTCATTTTGTACGCGTCACCCATCTTGCAAGA CAATAGACATGAAGTTATCGCAATTTGGGCACTAATAATTACTCATTTTCATC

>Gm15_899468	TACATTTACACGCTTATTTTCTCTCTTTTTTTGGATTATAAATTTCACTCACTTTATTACAGCTAACATCACTCAAATTGTAAACAGTCAAAAAACATAGGGTACAAATGCTAAGAAGCTAAGGTATGAATGCATTATGCATAGCCTCAAATGCATCTCTATTCAATAATTTAC[G/A]CCTTCTTGATTGTCTTGATGGATGACCTCAAGAACTGTTTGATGCTGCAAGATCGGCATCATCACTTGCCGTTGGCTTCTTCCAAATGTAGTCCACACCTTTTTTGATCCCAAGAAGCAAAGATGGAGATGATGATGGAATTCGTCTCAAGCTCCTAGCACTCTCTGATCTAGGAAGTCTT
>Gm15_3694443	TGACGATTCTGGCAGAATATCAAAAGAATTAATAATTAACCACACCCTAAAATGGCACTTATCTTGGTCTTGTTCTGTAGAGAACGTAAGACAAATTAATAAGCAACCCATCGCGATATTCAATGCTTGTCAACAACAAAAAGAATGAGTTGGAGGATGAAGTGATTTAAACAAA[C/T]ACCTCCTTCTGTTTAGCCTCAATAAGGGCAGCTTCTTTTTCTTTTTCAAGTTGAGCTGCAACTTCATCATTAAGTCTCTTCCGTCCCTCTTCCAACCGCTTTCAATCTCCACCTGAACCTCCTCAGATTCAAGCTTTCTTCAACTCTCTACGTATTGCTTCTTCAACTCTCTTTGCA
>Gm15_4176388	ATCTATAGATTTCTTACAATTTGTGGGGGATGCCATGTGGGATGGACCTTGCAATAGTAGCCATATATAGTTTCACTAGCTGTATTCTGGACTGCTGCCAATTGAGTGACTTGTTTCGATGTTTGCTTATTGCAAATCATGTTTATAGCTCACTGGCAATGTTTCTTTTTGTTAAGC[A/G]CAGGTACTCATATAATGAGGAGACTAGCGCATATGAAGAAGTTCACGAGGTTGCATCCAATGTGAGTTGAGTGACTAGTTTAAGGGGAGAATAGGCTATTTAGCATAGCCACAATCACTGTCTCATGTGGCTCATGTACAGACCGAAGCACTCTTGATCTTTTTGTACGTTATGGGGGATT
>Gm15_4176389	TCTATAGATTTCTTACAATTTGTGGGGGATGCCATGTGGGATGGACCTTGCAATAGTAGCCATATATAGTTTCACTAGCTGTATTCTGGACTGCTGCCAATTGAGTGACTTGTTTCGATGTTTGCTTATTGCAAATCATGTTTATAGCTCACTGGCAATGTTTCTTTTTGTTAAGC[A/G]CAGGTACTCATATAATGAGGAGACTAGCGCATATGAAGAAGTTCACGAGGTTGCATCCAATGTGAGTTGAGTGACTAGTTTAAGGGGAGAATAGGCTATTTAGCATAGCCACAATCACTGTCTCAGTGGCTCATGTACAGACCGAAGCACTCTTGATCTTTTTGTACGTTATGGGGGATT
>Gm15_6364520	AATCATTCTCCATGAAGTGTAGTGGTGTAAAGTGGATTGGAAAACTGGGGTGCAAAACAAGTCTGGATCATTCTCCACAGATGACAACAACTAGAGACTTCATGATGAGCCGATTCTTGCTGCTGCAAGGCTATGACTCTACTGCCTCCTCAATATTCTTCAAAAAAGCAATCT[G/T]CACTAGTAGAGCAACAGCCTAGAGATGTTAGCAAATTGGTTTCGCGATGATAAGAAGCAATTGCTATTAAGCATAACACTGCTATTATACCATACTGGCCAAAATCAAGAGGAAGAAAGTGAAATGAAGTTGATGATAATGAGTATGACAACTCTTCTAATATCGCAGCTAAAGGCT
>Gm15_39177610	AAAGAAGCTTAGAGGACGAGTTCGAGTTATGGCATTAGGTGCCTCACATTTATCACCCGATTTTATGGAATTTTGAAGATTTGCCTTGGTTGTCCGATGACTGAAGGATATGCAATGACCGAGAGTCTTGTTATAAGTTGCATTGATGAGGATGACAACCTTGTGGCTCTCCTAA[T/A]GTAGCCTGTGATATAAACTTGTGGATGTTCCAGAAATGAAGTATACATCTGATGATCAGCCCAATCCCCGTGGGGAATTTGTGTTAGGGTCCCATTGATGAAGATGGATGGCTAAATACTGGAGATATTGGGACTCAAGGTTGTTTTCCAGCGGAGGGTGCTGATAGCAATACAG
>Gm16_3592639	TAGCAAGAATGAAGAAATGTGAAGAGAAAAGGGAGCTACTCAGATCTGGGGCTGAAAAACGGCTATAAATAACCCGTAAACCACTCCTAATACCTCGCACTCTCTTCTCTGCTTCAACTTGCCTCCTCAGAAGCGAAGAAGAAGCCACAGAGAACTAGTCTCCTACTCTCACCCGCAAG[C/T]TACGAGACTCTCTCACCGCTTCTCTCCCTCTCTCTCTCTCTCTCACTTCTCTCTCTTTGTTTTCTTAAGTTTACGATCCGTTCCACAATTCTCATTTTCCCTTACTCTTCTTCTCTTCAACACTTCAATTTCACTCCGTAATTTTCTCAGATCTACACTTAAACGTAA
>Gm16_24688446	CTTGTGTTATGTTTCTTGCCAGTACTATTTCTGTGAAATATGTTCTCATTAGTGGCGTAAATTTGTTTCAGTATGTGATGTTCTATGCATGTGCACTGGACCCTGAAGGATGTGGTGTGAAATTTGCCATGGTTCTTGCTGATATGTTTGGATGTGATGTTAATCCCCCTATTACTAG[G/T]TAAATTTTTATTGTTGCATTCAAGAATTATAATTCTACTGAATCCTGATCGATAATGCTAAATAAAATATATAGTTGGAATGGGGCTTATTTATACACTCGTCACACTTGGCATAATTAATTTACTGCCATGATTTTTTTTCCATCTGATCAGTGTTGTTAATGGTGGATGGCGGTTCA
>Gm16_24688447	TTGTGTTATGTTTCTTGCCAGTACTATTTCTGTGAAATATGTTCTCATTAGTGGCGTAAATTTGTTTCAGTATGTGATGTTCTATGCATGTGCACTGGACCCTGAAGGATGTGGTGTGAAATTTGCCATGGTTCTTGCTGATATGTTTGGATGTGATGTTAATCCCCCTATTACTAGG[T/C]AAATTTTATTGTTGCATTCAAGAATTATAATTCTACTGAATCCTGATCGATAATGCTAAATAAAATATATAGTTGGAATGGGGCTTATTTATACACTCGTCACACTTGGCATAATTAATTTACTGCCTGCATTTTTTTTTCCATCTGATCAGTGTTGTTAATGGTGGATGGCGGTTTCA
>Gm16_35581397	AACTGAGTGCCCTTGCTGAGTTAAAGGCTGTAATCTCCATTTCTTCTCATCTTAAAGATCTCAAAAAAGATTTGGACATTTGTTGAAGCCTATTTATAGTCATTATACCCGAGTTGGAAAAATCAATGTGAAATAGTACGAGATGATGCATTGTAGAGACTAGAGAGCAGTCTTAAATTT[G/A]GAC

	TAGTTTACCCAATTCAGCCTTGTGGCCTGATAGCCATGTACACCACTTAGGTAAGCAATTA TAATGTACACTAAAAGTCTAAAAAGAACAATGAAGTACAGTACAGTAGCATTAGCATCCTTC AACTTTTGTATCTCAGTGTGTTTATAACTATTTTTTCAAGGACTGATTCCGT
>Gm16_ 3558950 8	GACCTTCATACTCTGAGCATCCTCATCAATTGTTTCTGGCATCTAAAGCTCACATCACTTTT GCTTTTTCTGAATTGGCCAACATTCTCGAGAGGGTTTATCACCCAGATGCAATAACCTCAC CACACTGCACCTTTTTCACGACGAGGGGGTAGCTCAAGGGTTCAGCTGAACAAAT[C/T]GAT TGTGTA AACAGCAGTGTGCATATAATCTTCTCTTCTTTTGTCAACAAGGTGAGCGCT TCACCAAGTAAGTCCGAGTCCATTGATCATAAGAGTGTATATCTGGACATTTAAATGGTAAT CTTATGCCGAGAGAACTTTTCAGCATCTTTTGATTAAATGTTACCATTTGAAT
>Gm16_ 3559056 8	CCTGACATCTTTACCTTTAGCATCCTCATGAAGGTTTGTGCCACCTTACTTAGATCAAACG GTGTTTGTCTATGCTGGGTAGTATTTTAAAGAGGGGTATCATCCTATCCATTAGGATCAC CACACTTGTGACGAATCTCTTTTTCTGTTGGAATGTTGAAGAAGAATTGAAGTT[A/G]GTGA TAGTGGGTAGCTCAAGGATCTCACATGATCCAGGTGGAATTGATCAGGTGGAATCCTTGG ATCATGTGATATCCTTGAGCTAATACACTATCATGAAGGCTTCTTTAACATTTCCAGGAAGA AAGAAGCCCTTCATGAGTGTGGTGATGGTAATGGTGATAACCCAGAAAAAAC
>Gm16_ 3559087 3	GAAAGAAGCCCTTCATGAGTGTGGTGATGGTAATGGTGATAACCCAGAAAAAACAGCATT GATCTGAGAAGGGTGACCGTAAGCATTTACGAGGATGAGTGAAGATATCAGGGCTAATCC CTTGTTGACTGTAATGGTTGAAAAGGGGAATCACAGTGGGATAGTGCTTCATCTTCAC[G/C] TGAGAAGTGAAGAACTGTTAAAGTGGATGATAGGAGGGGGAGATCGCTTAAAAAGCATG CGGTTGAACAAGGCAACAACATGATGGTTATCAAGGTGATTGTCAGTTTAAGACTGAGAAT GAGAATGAAGAAAAACGTGGAACACAGAGAGGGAGGGAAGAAACCTGAAGAAGGAAGAC
>Gm16_ 3559087 6	AGAAGCCCTTCATGAGTGTGGTGATGGTAATGGTGATAACCCAGAAAAAACAGCATTGAT CTGAGAAGGGTGACCGTAAGCATTTACGAGGATGAGTGAAGATATCAGGGCTAATCCCTTG TTGACTGTAATGGTTGAAAAGGGGAATCACAGTGGGATAGTGCTTCATCTTCACGT[G/T]G AAGTGAAGAACTGTTAAAGTGGATGATAGGAGGGGGAGATCGCTTAAAAAGCATGCGGT TGAACAAGGCAACAACATGATGGTTATCAAGGTGATTGTCAGTTTAAGACTGAGAATGAGA ATGAAGAAAAACGTGGAACACAGAGAGGGAGGGAAGAAACCTGAAGAAGGAAGACACT
>Gm16_ 3559105 4	GAGAAGTGAAGAACTGTTAAAGTGGATGATAGGAGGGGGAGATCGCTTAAAAAGCATGC GGTTGAACAAGGCAACAACATGATGGTTATCAAGGTGATTGTCAGTTTAAGACTGAGAATG AGAATGAAGAAAAACGTGGAACACAGAGAGGGAGGGAAGAAACCTGAAGAAGGAAGAC[T/ A]CTGTGACGGGCCAGGCCGGAATGGAAGAGTGTGGTGTGGGCTTTATGGCAAAAGGT ATGCAGTAAACAAAAGGAGGGTGTGGTGGGGTTTATCAACTTGAATGATGTGGGTGATG GAACTATGGAAGGTCAGAACCTGATACCTGCTTTCTTTTGAAGATAATCAACTTTTAT
>Gm16_ 3559117 1	AATGAGAATGAAGAAAAACGTGGAACACAGAGAGGGAGGGAAGAAACCTGAAGAAGGAAG ACACTGTGACGGGCCAGGCCGGAATGGAAGAGTGTGGTGTGGGCTTTATGGCAAAAGG TATGCAGTAAACAAAAGGAGGGTGTGGTGGGGTTTATCAACTTGAATGATGTGGGTGA[T/ G]GGAAGTATGGAAGGTCAGAACCTGATACCTGCTTTCTTTTGAAGATAATCAACTTTT ATAAATTTGTGATAAAAATAACAATAATAAATTCTTTAACAATAATTATATTACTAGAGATTGTA ATTAGATATTATTATTCAGGTTTAAATAATAACCCGCCATTTGATTGATGTTTTT
>Gm17_ 2714446	GTTTCCTTTTATAATAACCTATGATTTTATTCCTTTTATTCTAAAAATTAATTAAGAATATAT GCATGAAATGAAAAAATAACACTCCCTTCCATTTCTCCAAACCTAACAAACCTGGAAGT CTAAAAGATCCATCTCTTGCTGCAATTAACATGAAAATAAAATTATCCCAGGG[C/T]TTACGG GCATGGAACATTATGCCAAGCAGTTACATTAGGACGGGCTTCTTCTCTCTTTCTTTCCCTT CCACTTCCATATACTTCTCAATTTCCCTGATCATGGCCTCCATCGCCACCTTGAGAGGCTTC CTGCTCCTCGGAGATGTTATGGCCGGCTGGGAACCACCATCGCCAGAAGC
>Gm17_ 3909243	GTTTACAGGCACAACCTGTTAATGTTTCTACAAGTTTTAAGCAGAGGGCAGTGCTTGGA TGAGATCTGGCATGCTGCTGGTGATAATGCTTCCGACATTGATTGGTATGCCAAGCGCACT GTCCTTGGAGGAATACTCAACAACTGAGATTATATGCTGACAGATAGCTCTCT[C/T]G TTTGTTTTTCTTGAAATTTCTTCCCTCCCTTTTAAAGACACAAGCATTTCATAGAATTTGCTT GATTCTCTTTCATCCATTACCATTTCTAGTGTGTTTTGTGCCTTTCTTTTATGATGATCATTC ATTTGCAAGTCAACTTCTGTTTCTATGATTGCAGTGACCATGTTTTCTTTT
>Gm17_ 1036605 3	AAATATCCATGCCTACACCTGCACTCATCACCAACATATTTTTTTCTCAATTTTCATCTCTA GTGGCCCCACATAGTAATAGCTTTTTCTCTTGGACTATTTCCCTCCCTTTATTTTTCTCA ATTTTCACAAGTAAAGCAATTTTGGAAAATGGGACCAAGATTTTAACTTTACT[T/C]TACCTT GGCATTCTTTGAACTAGGCCTTACCAAAGTGGTTGAAAGCAATAGTGACCTGCATGCTCT TGTCTGAGTGTAGGAATCACTGTGGCCAAAAGCATGACCTTGTATGGTGAGTCATGTA ATTGTTGGAGATGGTAATCCCAGTAGACCATGAATGGCATCGATCAACCCA

>Gm17_11363185	AGAAATTAATAAACTTATCTGAAAAAATAATAAAACATTCTTTAGTCAGATTATTGGAAATGTTATTCATGAGATTTTGGACAGGTGATGAAAGTCTATCAAATGAAATCTCTGTTACTATGATCTATCACTGCATAAATGAAGCTTGATTAATTTGGGGATGTTGAGAGGCGCTCTT[G/A]CCTGTGGTGGCCCTGAACCCGGAGCTGGAGCTGACCCTGAAAATTCGGAAGAGAAAAAACAAACAGAGCAAAATTAATTAATATTTAACAATGTGCACTCAAAACAGGGGATAATCATCACGAAACAACCATAGAAAAGGAATATAATAAGATGGGGTTTGTACCGTTTGTGCAGCTGG
>Gm17_32028579	AATAGGGTGATGAGTGGGTGCGTAGACATCGAACAAAGATGAAGCTGAAGGTATATGCGGACCGCATGTCCCAACCATCCCGTGCGATTCTCATATTTACAAGTATTTTCTCTTATTATCTTGCTTTTCCTTTCTTATTTTATTTTGGAAATTTGAATTTGGTCTCTGCCGCATGTT[C/T]GAGTGAATGGAATAGATTTTGGAGTAGTCAAAGTCGATTTATCCCAACATCACCAGCTATCTCCCCAATCCGAGGTACTTAATTCGTTATTTCTTCTCTCTTTTTTTTTTTTTTTTACTTTCACGGAAAACGCTTTGTCTCTCTTACCCTTTAACTTTTATTGGATTTTCCCTTCA
>Gm17_39520782	TAAAAAAAATTAATTTTGTTCACCTTAGATCAACAACATAAGTTATTAGTGGTGACAAGCGATTAACTTGATCATGGTAGTAATAGCTAATAAATTTGATTTTAACTTCTTCTTCTCTTTCACCTACTCTTAACTTTTGTACTATAAATAACCCCCACTCACTGGCTCACTCTC[A/G]TACAACGCATCCTAAATAAGCATCTTAATTCATAGTCTCTTGGTCCTTCCCTTCTTCTGCTCAATAAGTGTGTGCCACTCTAATTAATTACCACCCCTTCTAGACATAGTTCTTCTCCCTCTGTTCTCTATTCTCTACACTGTGAAACCAAGATGAAGGTAGCATTTCAGCTGTT
>Gm17_41489915	GATGATAAACGGCACCTCGCAGCACATGCTCCGTTACAATCTGAAATTAATTATTACCATCAACACATAAAAAATTTGTACTTAACTTCCAATTTCAAATAGTTATGAATTTATATATGAAACTTTAGTTAGACCTATCTGCTGAAGAAGAGAACCCCTGCGTTTGTGCATGTGC[G/C]TGCGGTTCCCTCAAAACATGGTCAAATTAGTTAATTTAATTTAGTAGAAGTGTAAAGAAATTAACAAATTCACCTAACTTGGTTACATAAAACGAACAAGAAAGAGGCTAACATATTTATATGAAAAATTAAGAAGCAGAATATTTGAATATTTATGTACCGATTGATCAGCA
>Gm18_1392264	AGGCTCCATTGTCCAAACACATTGTCTTTTGTGTATAATAACTGGCATGCATGTCAAATAAGAGAAGCAAGCTAGAAGGGGTGAAGAATTGAAGCACCTTCTCTGACAACACTAGGTTCCCTAAATGTGTTGGAGCTCTAGTAACTCAGAGGTGGTTGCTTCAAACAAAACCACGTTA[C/T]AAGAGGTTTTGGACATTGAGAACAACATGGCTCATCAGTGGGTGCTCAATGCTCCAGAGCCACCTTCCATGCTGCGTCAGGTGGTGGACAACGTCAAGGAAACCCTCTTACCGCATCCAAATCCAAACACATTCTCTTATCTCAGGAACCAACCCTTTTCCAAACGTGCCTTTGCCTTGT
>Gm18_1392269	CCATTGTCCAAACACATTGTCTTTTGTGTATAATAACTGGCATGCATGTCAAATAGAGAAAGCAAGCTAGAAGGGGTGAAGAATTGAAGCACCTTCTCTGACAACACTAGGTTCCCTAAATGTGTTGGAGCTCTAGTAACTCAGAGGTGGTTGCTTCAAACAAAACCACGTTA[G/T]GTTTTGGACATTGAGAACAACATGGCTCATCAGTGGGTGCTCAATGCTCCAGAGCCACCTTCCATGCTGCGTCAGGTGGTGGACAACGTCAAGGAAACCCTCTTACCGCATCCAAATCCAAACAATTCTCTTATCTCAGGAACCAACCCTTTTCCAAACGTGCCTTTGCCTTGTGCAA
>Gm18_2151360	TACTAACATAATCAAAGAGAAGTCATGTTCAAGTGTGTTAGTATGTAGTGGTTTAGATAAGTTCATGATTGGGCAGGTACAAAGTGTGAGGTCTGATGGAAGCAATTATTGGATCAGCAATGAGGCACATGTTGGTGGAAGTGTGATGCAGATGATGCTTAGTCATCTACCATTCCCAAT[T/C]GTACATTCCCTAACCCCTGTGTTGTTCAATGAATGTGGGTTTATATAGGAAATTTATCAAAAACAAACGTAGTTGAACATATTGCCAAACCCTGTCAATAACGTATTTAAGTACATCTAAATTTTTATCCTAATGTTTTCTTTTAAAGAAAAAACTTATTCAAACTTCTAATTTGTCTGT
>Gm18_2235005	TTTTAAATATTTTAGCTGTTAATTAATTATGCAATAGAGTCACACATTTGGATTGTGCTGAAAGGATCTAGTCAATTTTTTATTCTATAATAATGGAACCAATGTTCTTTTGAATGATTGTAGCTGATTTTTTTGTTTATTCTGGTAATGGCAGTTGATATGCTGAAGGCAA[G/A]TGCAGTTGACAAAATTCCTCAGATTTAGAAGGGATGGAAGCATCAATGCAACATCTGCTAGTCTTGA TTGATGACATCTACAAATATGTTGACGATGTTGTGGTACGATATTGTTGCCTTCTTTAATCTCTCGCCTTGTCTCTGAGATTGTATTATTATATAATTTTTCAACACAGGA
>Gm18_2235806	AAACGTGGAATGTCAGCGTTTTTACTTCACTTTTACTTCGCTTATTTTTTATTTATGTTTTGTGCTGGTCTTTGCTAGGTAAGTTCACCTATTGTCCAAGGTATGCAGTGTAGCGTTCACCTGTATTTTTTCATGTAATCTGTATTTGAAAGAAAAATGAGCAAAAAGCTTTCTCTCC[C/T]TATCTATTTATTTCTTATTTATTGAGATTGGATCAAGTTTCAGAAAAATTAACAGTTTCTTTAGAGGTTAATTAGTCATTTTTTTATTAATTTCTGAAATTTGAGACTTTTTCTTAACAAATTTTGACTTCGACTTTTCATGTATGAAGATTGTTAGATTGATAATCAATTAGAAAA
>Gm18_8379461	TCATTAGTGATGATGTGATATTAATATTATTTATTATTTTCTCGTGCCTAAAAATGTCAAAAATTTTGCTTTAAATGATTTAATACTGATGCATGATAAATAACTTCAGACAAACAAGTTTACGTGAAAAATTATATCAGGCAAAAAGCTTAATGCACGTACAAAATCAATCTATACGIA/GIATTACCT

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>Gm18_ 8828934	CTCAACTAAGCTTCAAGATCACATTCACAGCTGATTAACTACAACCAAGAAGGGTAAAAGAA CTAAGCAACTAAAAGAAGAACTTTCTGTTGGGGATGCTCTAGATGCAACCGTTGATGGATAGT TTGAGGACTTGAGAGGAGGAGACACTGCCTTGCTTTCTTGACCCCAAGCTCTCTG[G/C]C AAGCAACTTAGCTGCCCTTGATCTGTTAGTAGTGTCTTCTGGAAGGACTGTGAGAAGCCA GTTGCAATGATAGTCACATGAATCTCCCCATTGTAGCGATCATCAACAACGGCACCAAATAT AATATTGGCAGAAGGGTCTGGCTAAACTTGTCAACCACTACAGCAGTTATGAAGTTT
>Gm18_ 8834781	ACTTTAAGAGGGGGAGACAGAGAGAGAGAAGAGAGATCATATATGGCTATTACCCAAACG AATGCACAACTAATTTATCAATTACCCCACTCAGCAGGCAAAATATTGCTTTTCAAAAACATC ACTATAATTGCAGATGTAGAAAAATAAAGCCACACCTGGCATCGCAATCACAGATT[G/T]TC CACTGCCTCTTGAAGTAACCTGAGTTGGAGGAGAGAAACAAATTGCCAAACGTTTGGACCTT TTCATGCACGCATTGTCTGTCTGCACCCTGAAAGGTTTATGACTTCAGTAATCAGTAAAGA GATGTTTCATTGAAACAAACCAATTTATGCATAATGAAACCTAGGTATTCAATAAA
>Gm18_ 8846329	AGTTATGGTAAGAGTGTTACCCTCAGCCGAGAATCTGTAACATCTACATGTTGGTCCAAGT CATCCTAACAAAAGAAGAGAAAGGATTAGATTTAGAAATGAGAGAATAAAATGGACAGTA AAAGTTGCAAAGAAAATTAACATATGAGGGGAACATACAATGAGTCTGGTATGTAGA[T/A]T CAGCTCTTCATTACAGCCAACGCAATATGTTTAGTACTGATTACAGTCTCCTCCAATTTCT CAAGGCCATCGTCTTGCTCTGCTTGAATAGCACATTTCCAAATTTAGATCCAAACAGCAGAC TATGTATGGGTAATATTTTTCAATTTCAAACAATTCAGATTAAGTGACCTTTTCAT
>Gm18_ 9028509	ATTTGGTACTGAATTTACTTGTCTTGTTCCTTTTAAACAAGGCAAGTAAACTATTCAAAA CACAAGGACTCGATACAAAACACGTGCATACATCATGCAATTGAATAATTGCACAACATGT ACAAGGTTGCATACAAAAGTGCAATGTACTAATATCTTAGTTGAGCTACACATT[T/G]TGA TCTCCTTTCTTTTACTCTTCATTTTAAAGGTCATCAGCATAAAAATAAAGACATATTATATCATG CAACTAAAAACATAACTGAGTCTTGACTCCTAACTAAAACTTGAGCCATCTCTTCTCTTTTTC TTTGCTTGTTACTGAATTTGAGGCAGAGGCATGAACTGCTTCAGCATC
>Gm18_ 9133223	GTAAAAATAGAAAAAATTAATAATTCCTTTCTTCAACGAACCAATACATCATTTCTGTGCC GTTTTCAATTTCTTCTCTTCAGAGGCAAGAACGGTGCAGAACAGATCATCGGTTGCCACG TCAGCATCGTCCAACGGTAACACGTTTAGATCAAACGGCAACGGTTCTCGCCGGA[C/G]CG ACGACGCCACGTTGTCGCCGTCTCCGTCTGCCACAACGGATGACGACGAGTCCGACGTGCG TGCGGCAGTCCCTCGGGAACGACGGGCGGCGTCCGGGGATATCTCCGGTTGACAATCACC GGCGCCTTCGCCGGCACGGCGGCGCGTGGGCCGCTGAAGGATTCTACGGTGCTGCTCAT
>Gm18_ 9259541	GCATCTATGGAAGTAACTTTCAACCAGAATTTTCATGACATTTTACCACTGAATGCTTTTCTTT GGTATGTCTACCAATGTAAGAAAAGGATAAGATTAAATATGATCCACAGCATACATGTAAT CTACATGAAGGTGCAGATGTTAATTGAACGGGTTGATGGAAAGCCACTTTCTAA[A/G]ATGT GGGTTATTCTAAAATCCCTGTGAAAGCTTTTCAGAGCAATGAATCCTAATCCTAAATTTATAC AGATGCAAGATACTGAAATATGGCACTTCATGTTTAAATTACATAAAACTAAGATCATGCTT GCAACCCCATTTGATTCATGTACCACAAATGTGCTAGGTTCTGTGTAGAGTT
>Gm18_ 9299254	TTCAATTTATGATAGTATTTTGGATATATAATATATATATATATATATATATATATATATA TATATATATATATATATATATATATATATATATATATATATATATATATATATATATATA CACAATCACAACCTCCCTCTATATCTCTCATCATCTTATCTCTCTCT[A/G]CTATAATACA AGACCCTAGTCCCTCTTCACGCCCAACCAACCATCAACCAAACCTTCCCTCTCTCAATTT CCCTTCAATTCCTTCTGGGCTTTGCAATTCGGCCAATTGGGTCCCCAAATCTGTAACCTTT TCCTCTGTCTAACGCGAAACGGAAGAGAAAACCCACATAGAGAAA
>Gm18_ 9299278	TATATAATATATATATATATATATATATATATATATATATATATATATATATATATATAT ATATATATATATATATATATATATATATATATATATATATATATATATATATATATATAT TCTCATCTCTTATCTCTCTCTCTCCTATAATACAAGACCCTAGTCTT[C/G]CTTCACGCC CCAACCACCACCATCAACCAAACCTTCCCTCTCTCAATTTCCCTTCAATTCCTTCTGGGCTT TGCAATTCGGCCAATTGGGTCCCCAAATCTGTAACCTTTTCTCTGTCTAACGCGAAACGG AAGAGAAACACCCACATAGAGAAAGAGAAAAAAAAGGAGAGGAAGAA
>Gm18_ 9299279	ATATAATATATATATATATATATATATATATATATATATATATATATATATATATATATA TATATATATATATATATATATATATATATATATATATATATATATATATATATATATATA CTCATCTCTTATCTCTCTCTCTCCTATAATACAAGACCCTAGTCTT[C/G]TTCACGCC CAACCACCACCATCAACCAAACCTTCCCTCTCTCAATTTCCCTTCAATTCCTTCTGGGCTTT GCAATTCGGCCAATTGGGTCCCCAAATCTGTAACCTTTTCTCTGTCTAACGCGAAACGG AGAGAAACACCCACATAGAGAAAGAGAAAAAAAAGGAGAGGAAGAA

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>Gm18_1092285 1	GCGTTCGGATTGAGCTTTCTTCAACCGCTTGGTTCGCGGAGACGGAGAGCGCAACAAGGGGAGAATCCGGCGAACCGGGCCGAACCGGTTTGGCTGTACTGCGGCCGGAATACTCGTGCTTGTCTCGCGCGCGGAGAGATACTACTGCGGTGGCGGCGTGAGGAGGGAGAATGGT[C/G]ACTGGAGTGACGGCCCATGGTAGTAACGGTGGAGGTGCTGTATCAGAATAATTGAA TGTTCCAGAAGTGTGGTTAAGTTACTAACATCACAATGTTAAGCATTGTGCATTGTTTGTTA CTGCTTATACGCTCTGTGATCGCGATTGATTGAGATTCTCTATTGTGTGGTGAGTTCAGAAT
>Gm18_1120957 2	TGTGGCCTCACTAAGGGATGAAGCATATAAGCAGCCTCAAGCACATTCAAGTCTTCAAGGT TCAAGCACAAAAATGGTAACCATTTGTGGGTTCTTACAATGTCACTCCTAATCAACCAACTCC AAAGGATCCTTCATGGCTATCCGATAGTGACCAAATCGGTGTCCTAGGTCATGTAGT[C/C]CA TTGTCTACATATACGAAGCAAATCCCAACAGCAACACCATTGAGAGATTGAGGAACTCTCTT AGCAAGCTTTTGGTGTACTACTATCCATTGCTGGCAGATTGAGCTTGACAAAAAGTGGTC GCATAGAAGTGGATTGTAACGCAAAGGGAGTGACATTGATTGAAGCTAAAACCTC
>Gm18_1120957 4	TGGCCTCACTAAGGGATGAAGCATATAAGCAGCCTCAAGCACATTCAAGTCTTCAAGGTTCAAGCACAAAAATGGTAACCATTTGTGGGTTCTTACAATGTCACTCCTAATCAACCAACTCCAA AGGATCCTTCATGGCTATCCGATAGTGACCAAATCGGTGTCCTAGGTCATGTAGCC[C/C]TT GTCTACATATACGAAGCAAATCCCAACAGCAACACCATTGAGAGATTGAGGAACTCTCTTA GCAAGCTTTTGGTGTACTACTATCCATTGCTGGCAGATTGAGCTTGACAAAAAGTGGTCG CATAGAAGTGGATTGTAACGCAAAGGGAGTGACATTGATTGAAGCTAAAACCTCAC
>Gm18_1193421 5	ATTAATCCAATTAAAAGTACAAACATCCAAGACATTTGTTTCAAGGTTGTGTCTTCACTGCAA ACACCTAAAATGGTTATGGATTGACTTACAAGTAGGGACTCAATTTTATTGTCAACAGTCT CTTTGATAGTTGTAACCAAAGTTTACATCTTTGTAGGCCAGAGAAAGCTGGCAC[C/G]GTTT CTAGGGAAGCACTGCCTCTAATAAGCCCTCACTCTCTCATTTTTATACAAACCATCACCAG AGACTACAGCAGCTCCATCCCTAAAAAGTTCCTTTTCTCTACAAGGATTGCTCCCCCTGG CCAAAATATTCACAAGAAAAACACCCTTCTAAAACATTCTTGTCATACCATAC
>Gm18_1194013 7	ATCTCACACAACATTCAAATTAACAAAACACACAAAAAAGCAATCATTATTTCCCAATCAAAA ATCACACAATTACAACGCCAATCAACACAAAACACATATACAAAACTCCACCTTTTTCTGTG ATCTTTGCTGCATTTTCCCCTGACACCCACAAATTCTCCAAAACCTCCCTATGGT[C/T]GTGAT CAAGCCCAAACTTAACAAAACCTAGTTCCCAAAATTCTACTCCTTCAATAAAAATAAAATGAAA CAAATATTTTACACCTTACTCTCACCCACATTTCCCCCACCTACCCCTGCTCATTATCAC AAACCCACAACAATCCCAGCAGCTTATTCTCCCTTCTTCCCCCAAACCTT
>Gm18_1194013 8	TCTCACACAACATTCAAATTAACAAAACACACAAAAAAGCAATCATTATTTCCCAATCAAAA TCACACAATTACAACGCCAATCAACACAAAACACATATACAAAACTCCACCTTTTTCTGTG ATCTTGCTGCATTTTCCCCTGACACCCACAAATTCTCCAAAACCTCCCTATGGG[A/G]TGATC AAGCCCAAACTTAACAAAACCTAGTTCCCAAAATTCTACTCCTTCAATAAAAATAAAATGAAAC AAATATTTTACACCTTACTCTCACCCACATTTCCCCCACCTACCCCTGCTCATTATCACAA ACCCACAACAATCCCAGCAGCTTATTCTCCCTTCTTCCCCCAAACCTT
>Gm18_1311316 1	TCCGTTAAATAAAGATCAGCATAATATATGCTTTTCGGCTATGTTGATCCGAACAAAAAGCA GCTGGATTTCAAACACCTTAAATACTGCCAACAATGAGCACAAATCATACAATTACAGAAGC CTTATTCTATTAAGTGAGGTGCGCTACATAGAGCCACGATGTCATTAGACTTAG[C/T]CAA AACCAATATTCATAGATACAGTCACCAATCATATCATGGCAAATTCAATAGTTTTCTTCCCA TCATGAGCCTTGCTGAAGAAATACTTGAAATAGTCAGAGTACAAAAGTTGTTTATATTTTGG TTCATCACCATCATCCAACACTTCCGACAAAGGACCAATAACAGCATCAGGT
>Gm18_1311350 0	GGACCAATAACAGCATCAGGTGCTGGGTTGACAAAGATTGGAATGGAGATCCTGGTCTTG CTTCTATTTGCAACCACTCTATGCTCTATGCTTTTGCATCTTTCATTGCTCATTATCTGTAGC ACATCCCCAATATTGATTACTAAAGCTCCCTCAACAGGAGGAACATAGATCCAACCT[G/A]TC ACCATCACTGCCTCTTACATAAAGCCCTCCAATATCATCCTGTAGAAGGACAGTAATGGAT GATACATCAGAATGAGGACCTACCCCTGCTACAACCTCAGGGTCAGGGCAAGCTGGGTAG TAATTGAAACCCAATATCATTGCCCCCATAGAGTGTTCTCTTCTGCTTTATCTAAC
>Gm18_1311350 1	GACCAATAACAGCATCAGGTGCTGGGTTGACAAAGATTGGAATGGAGATCCTGGTCTTGCT TCTATTTGCAACCACTCTATGCTCTATGCTTTTGCATCTTTCATTGCTCATTATCTGTAGCAC ATCCCCAATATTGATTACTAAAGCTCCCTCAACAGGAGGAACATAGATCCAACCTA[A/G]CAC CATCACTGCCTCTTACATAAAGCCCTCCAATATCATCCTGTAGAAGGACAGTAATGGATGAT ACATCAGAATGAGGACCTACCCCTGCTACAACCTCAGGGTCAGGGCAAGCTGGGTAGTAA TTGAAACCCAATATCATTGCCCCCATAGAGTGTTCTCTTCTGCTTTATCTAACT



>Gm18_1320870 5	TTTAAATTTTAAACATCTATCATGAAAGATAACATGATCCAAGGATATGCATTGAGGATCCAA AAGGTCATATCCAAAAATTTAGTTCATGTAAATCTCAACATTGAACTACTTCAATTTTTAACG GAAAAAAAAAAGGAGAAAAACATTACTAAACTCCCAATGCCATACTATTTTC[G/C]CAATAA TTCCCATCACTTATATACAACACAAGCGGTGTGACACACAGCTTGATAGTTCTATATTACCA CAAAATTTGAATAATCCACTGTTTGTTCACCAATCAGCATTTTGAACGAGGTGGATTGAG TTTCCACAAAATGCCACAATATCTCTTTAACCCAGACAAAAGAGCAGAGAT
>Gm18_1320993 6	ACTCTCCTCAAATTTCTTCAGAGAGACATCATCACCAGGGTTGTTAATTGGATACACAACT GGGACTTAAGTCGAGGCTTACCATTGGCAAGTCTTTTGTTCAGGGTAACTGTTCCAGG AGTTTGATCTCGTCAGATAGCTTCATCGGGGTAGGATTAGGCGATGGTTTATATG[T/C]AC TCTCATTATTTGGTGAATCTGTTTCTTCATATGTCGTCGCCATAATCAGATGAATCACAG GATGATAAATCACTGCCACATTCAAACGTACAACTTGTTCCTCCTCTTAGTGTATGTGGT CGATGGCCAAGGTGAAACAGAAGCTGTATTCTTGCCAACTGTGAAGACTCCC
>Gm18_1342655 7	TCTGACAACACAGTAGAGAAAAACAAGAAGTTGATGCCACCTTGCTGAAGCTAACACCAG AACAAGTGGGGAAGCTGAAGAAGAAGGCCAATGATGACTCAACAAAAGAAGGGTCAAGAC CTTATAGCAGATTTGAAGCTATTGCTGCACATATATGGAGATGTGCCTCTAAGGCTCG[C/T] AACTAGATAAGAATCAACCAACTCTAGTTTCGGTTCAATGCCGATATTCGCAATAGACTAAT TCCACCTCTCCCTAAGAATTATTTTGGGAATGCTTTGTCCTTAACAACAGCATCATGTCACG TTGGAGATGTCATATCAAATCTTTGAGTTATGCTGCTCAGAAGATAAGGGAAGCA
>Gm18_2103071 2	TGGATATAGCTCTCAGGGTTCTGGATCCACACTTATTATGCCATTTCTTGATAACCAGCTGA AGTCTCCTAGCCCACTCCTGTTACCTGCGCAGGTGAGGTTCTGAAATGGGTGTTTTGCTTG AAATATTTGATTCTTCAAGTTTATTGACTGACACCTACATTAATAATTATGATTTCT[G/A]TCAG GATGCTGTTACTCCATTGTCTGAAGCAGAAGCAGTTGATCTGGTCAAACTGTTTTGCATC TGCAACTGAGAGAGATATACACTGTAAGTTTGTGCTAAATACATAATTACCAATTGACTG TAGGATGAACCTGTAACCTACTGTGGATGCTTTTACAGGGAGATAAGGTTGA
>Gm18_2483025 1	TTTGAAAATCGAGCTTAGTGCATATTTTCATGTTTATCTTGTTTCAATCTTCCCTAACTGCC TCTTGAGGTTTAGATGTCAAAGTTTCTAAGTATCCTTTTAAACAGCTGAGCTCCAGTTCCAGAA GAGAAGACTGAGGTGGACGCATTCAATTCCTCATTAGCTTCACTAATTTTCATCA[C/T]GTCC GATTCATGTCAAATTCGTAACACACACAAATCATGTGTGTGTGCTCATTACTCAATTTT AAATTGCCATCTGAAATTCGAATTGTGTACAAAATGTTGTGGCTGCATTGCTGAGATTTT AGCAATCACAAACACAATTGCAACTGCAATCACAATCTAAGCCATGCGCATT
>Gm18_4876691 0	CAATTTACAATATTTATCACTGAGCTCGGTGATGGTATATTGACTAATGTATTGTGTGGCTC TGTCTATTTGTTGAAGACTCTAAACATCCTACGGTGGGGGTTGCCTTAAACAAGCATTGCTCT TTTGTATCGGCGTAAAGCAATCCAGGAACATTCAAGTTCGCTGTTAGTTCAAGAG[A/C]TTG AACTCTCCAATCGCTGGTTTCTTAACATGGAATTTGATCAACTTTAACTTTAGTTCTCATT CTCAATCATAGGTGCTATGAACCTGATATATTTTGTGCTATATGCAAAACAAATCAAACCTATT TTTATTGTAATTCAAACAATTTCTTATAATATAAGAATGATTGACAGGGTCT
>Gm18_5301517 8	CCCCGTAGGTATCCAGCTTCAAACATTTGATGCATCTTCCTTTGTTGGTAATGCAGAATTGT GTGGAGCACCTTTCCTAAAACTGCAGTAATGAAACATATGGGCTGCCATGCACATTTGG TTGGAATATCATTATGGTTGAATTGGGATTTGTTTTGGCCTTGCACTTGTTATTG[G/T]TCC CCTTCTGTTTTGGAAGCAATGGAGGCAATGGTACTGGAACGCGTGGACCTCATTCTGTGT CGTATCTTCCACAGCTGAATCTTGAATATGAAAGCAGTGGAGGACATTGCTATCAAGTTTT AAGGTGGAGGCATTAGTCATGAGGGAGTACTAAAATGCTAATAGCAAATCAGCC
>Gm18_6054804 5	GGGAATGAAGATATAATCCGTTTTAAAAAGGTTGACTTTTCTTCCCCAAGTGCTTACACGCT GAAATGCTGCTTTACCATCACAAGCAATTGATGTATCATTTCTGCTTAAACCAGCAAGCTTA GCAAAGAAAAAGCAATTGAAAAAGTGAAACCTGTCTTGTATCGCTTGGAAGTT[T/A]ATTT TTCAAATAATTGATATAAAAGTCAATGATAGTATCATTAAAGCATTCTCAGGCTGTAAAAGC TCAATATCGCTCTTTTTGATACAAACAGCATCAGGGTCCCCCTTTGGATACGCCTTTTTTAAA TGCAGCAGAATCATGCTCCACTATCTGGTCTCCCAAGTTAATTTTCATCTTC
>Gm18_6054805 7	ATAATCCGTTTTAAAAAGGTTGACTTTTCTTCCCCAAGTGCTTACACGCTGAAATGCTGCTT TACCATCACAAGCAATTGATGTATCATTTCTGCTTAAACCAGCAAGCTTAGCAAAGAAAAAG CAATTGAAAAAGTGAAACCTGTCTTGTATCGCTTGGAAGTTTATTTTCAAAT[A/G]ATTGA TATAAAAGTCAATGATAGTATCATTAAAGGCATTTCTCAGGCTGTAAAAGCTCAATATCGCTC TTTTTGATACAAACAGCATCAGGGTCCCCCTTTGGATACGCCTTTTTTAAATGCAGCAGAATC ATGCTCCACTATCTGGTCTCCCAAGTTAATTTTCATCTTCTGAGGGATTGAA
>Gm18_6071144 3	GCACTGTCACTCACATCATAATTATGAAGCTGGCAATATTGATTCAATATTTCAAGGGAAAT AAAATAAGGAGATGGAGTTCTCAAACCTTATCAAGTTCATTGAGAAGATACACACAAGGGGG GAATGGAGGGGAGAAGAGAGAAAGCGAGGTACTTGGAAGATTGGCTTAAGGTCAA[C/T]G

	GTCAACTGAACAGCAATAAGGGAATCCTCATGATCTCGGGTACCCAATACAGCTCTCGAGT CACCAACATTTCCAATAACAAGGTTACAGCCCTGAAAGAACATCAGTAAACCACATGTAAAA AGCAAATAAGAAATAATCAAGAGAACATTACATTGTAATTTATATTTCTTACCTGC
>Gm19_ 451849	TGAAGAGGACGTGATTGCCCCCAAACAGGGACTCAACTCCACAGGGCTACAAGTCACAGC CTCTGCCATGGGACTCATCTCCACCAGAAAAAGTGCTGCCATCACCCTATTGCAAACCCA CACACCTTCTTCATCATATGCATGCTTTCTTGATTTTATTTAATTCTCTACTGAGTGC[C/T]AT TTTCTATGTGCTCAAACATATGCTTTTATATACAAACAATTCAAACACGGTCTGTGGGGTGGGA CTCAAATCTAATTAACATCTAAGGTCCCACTTGTTGTGCTAAAAACATAATGACAATAAGTGA TAAACCCCTGGCTGTGTTTCACACTCAATCTTTGCTGAAATGGTAATACGGAAAG
>Gm19_ 829824	ACAACCTGCGACTTAGTTCATCCCTTCTCAAGCCACCGAAGCCTCACTTTTGTATTCTTCCCT CTCCCAATATTTTCTTCTTCTTCTTATTATTATTATTCATTTTTTCTATATAAACCTCAC CTTAATTACTCTACTCTTCTCTCTAATTACAATTTTCTATTTTGTTTAATT[G/T]GTGAAATTA CTTGATAGATAGATAATCATGGCTATGATGAAACGTGCTGCTTCCCTCTGCGCTTCGCTCTC TTATTGCCTCATCCTCCACCTTACCAGAAACCTTCATGTAAGCCCAATCAAATGTTACAAA TTAATTGCATGTTTTTTTTTAAAAACATTATTATTTTCTCAAACCTCA
>Gm19_ 1567798	TTGATTTTACAATTGCAGCACAACCACACTACAATTTAAAAACATTAGTGAAAAAAGAGGGCA TATCTATATCATTATCAAGAGAAGATGAACATAGAAAATGAAAGATTATGCCTATAATATTCT AAATAAAGCAGTAGAGAGCAAAAAATTGGAAAACCAATATGAAAAAAGCAGGC[T/C]TACC TTGGATGAGCTGCAAGCTATCTTACATTCCAATCCGTTAACTACTTTAACACCTTCCATGGC AGAGCATATTGCAGAAAGCTCATCTAAGTTTCCATATTGGCGTTTGTTCGATATATCAA AATAAAAAGTAGAAGAACTCTGTTACCTAGTTGAAAAGGATATGCAATATT
>Gm19_ 1992839 5	CACAAATGAAAATACCTTCAGATCATGAATTAGTTGGGCAAGATCTTCAATTGAATAAATAT CATGATGGGGAGGTGGACTGATAAGTCCTACACCGGAAGTTGAATTCCTAGTGACAGCAAT GTCTCCTATAACTTTGTGACCAGGAAGTTCACCTCCCTCACCTGGTTTTGCTCCCT[G/A]AA GATGACAATTATAATGTCAAGACACTACCATTGGGTAAGTGAAAAACTATAGCTTTTTATTAA CCTTACCAAAGGTCAAATGCAGGTATACTTACGATACAACACAAGAGCACCCAGAGGTGCC TTGAAATCATATAGCATCATCTTCCCTTGAATATAAATTCATAAACAATAGAAATT
>Gm19_ 3493361 7	TGAACCATCACTCCACTTGGAGACCCTACTTCTGTACCTAAAACGCGCACAAAGAGAACA AAAAGTAACCAACAGACACCCACACACAAAGACACAAACACGCAATTCTACTATTCAACACT TCTATTAAGGTCATAAGCTTCCAGAACTAAGGAACACAATATAAAATACTACCTT[C/T]CCG GAAGATCCACCAGAACTTGAGCGGCTGAAGACGACGAACCTACCGACAAAATCGAATCA CTAAATCAGTTAATCAAGCAGAGCACCAAACAACGAAAAACCATTTAAACACACAATCTCAA ATTTAGACAATTAATCCCCAAAACAAAATAAAGCAATTAACACCACAATTTTCG
>Gm19_ 3685686 1	ATGGCGTGACGGTCTTGCTCGTGCAGACGATGAAGGTCCTGGCTATGTATTGCCAAATAGA ATTATTTCTTGAATCGGTAAATGAAATTTATCTAGACTTCTATATTGTCTTTTGTCTGATAAGT AGAAGTTTTGCACAGCAAACTACTTCACTTTTCTTTTACCCATTTC[A/G]AGCTAA CCAGATGCCTGTCAATTATAAGCAATCTACGGCGAATAGTAGGAAGGACAAAACACCCATAT GTTGAGCGCAGCCTTGATGTCAATTGTCAACATTGTGACGACATTCTATGCAAAATGCTGCTG CCTTTGAAGAGGCTGCACTACGTTTGAAGAAGAATGCTGCAAGTGTAAG
>Gm19_ 4342540 7	AAAAATGTGATAAAAAATAGTACATTCCTAACATTTCTCTTGTATCATAAGACCCACCTTGG GTCCTAGCATGATGCAGGGATATCTAAGAGAACTTCTTTCAAGAGATCAGGATTTGTTCA TATCTATCTTGCAACTATATTTAAGCAGAACAGAGTATCATAGCAAATCTTGTT[G/C]TCGCC ATTTTAGTAATGTAATTCACCAAGTAAACATAATTTTCAATCATCACTACAACAAGCTTTGG AATTCTCCATGGGACCAATTCCCTAACTACAAAACAAACAGTCCCCACTGCTACTACAAAAA ATTTAATTTCCATCATCACCAATTTCCCTAATTGCTCTCTTAGCTGAGCAT
>Gm19_ 4703883 6	TCAGCAGCATTATAGCATTTGGAAGATCTTGCTTGTGTTGCAAAAACCTAGAAGCACAGCATC TCTCAACTCATCCTGACAAAAATTATAGCGACCCATACAAGAGAAATAACAAATTAAGCTA GGAAAACGGGTCCTCTAACTAACTAAGCCAATTTCTCATATTGAATCGTTTAC[G/T]ACCT CATTCAACATCCTGTGCAGCTCATCCCTAGCTTCAACCACACGATCCCGGTCAATTGCTATC AACCACAAAGATAAGCCCTTGGGTATTTTGAAGTAGTGTCTCCACAGAGGACGGATCTAA TAGGAAAAAGTTAGAAAACATTGCGGTGTTCACTACTGTATCTCAAAAATTTAAT
>Gm19_ 5041340 6	CCTCACTCTCTTGCCATTTTCTTTCTTTTCTTTTGCAGGCCCGCAAAGTTATTGCAATGTT GAGTCAGGATTTTATATTTGATGGCTGTACAATTTTGGTTCATGGTTTTTCCAGAGTCGTCT TTGAAGTCCTGAACTTGCTGCACAAAATAACAACTATTTTCGGGTCTTCTGCA[T/A]TGGTT TTTGCCCTCTTACTCTCATTATTTATATGTAACACAAGCCTCATTTTCATCTTTAGACTGTCT TGGCAAAAGCAAACTAGTTTCATAACCCAGCTGAAAATTAAGTTAATTGAAAGCTTATA ATCTATTATAAATTAGGACCTAAGCAGCTGATTGAATTACATGTAAATTG

>Gm20_12115300	TGTTTTTGTGGCTATTATACAATATACATATTGTTAGCCAATCTTAACTGTTGTAAGGGCTTAATCTATAAGGTGTACCATTTGAGATGCATTGTGGTGGAATATTGATTATATGCCATATCTAGCATCTTTTCTATCCTTGCCTATCATGTAATACGACTTTTCTTTTCTCAGGAGCT[C/T]ACAAGTCAGTTTATGCCCCGAACCTTTTGATGTTGGACGCGTATTGCAGGTTGATATTATTTAGAGGGCCACCAGATCACATTGTCAACCACTGGTCCAATTGATCCAGGTTGGGTGCCATGCCAAGTTGTACACTTTTAATTTAGTATTCATTCACAAGTATCATTCTATGTATT
>Gm20_25068927	AACAAAATTGCATAATCATATGCCTCAATTGGACACAAATGACCTATCCAAAATTTTGTTCGTGACATCCCAAAATAATATATACATAATGGATCAACAGCAGCAATTTAAAATCTTCGAAACAAAAATTGAAGTTAAGAGGAACCCCTTATATCATGTTTAAATAATGGCTGTGTTGGC[G/T]GTTCATTCATTTCCCCCATAAACTATGGCAAGTCCATTTACCCCATCACTCATTTCATAGGCACCTCAGAGCTGACTTGTTTCATCGTCATCCTCACTCTTGCTCTCTTCAGCATCCATTGTCGCTCCCTTTTGAAGTAATCAACTTTGAAGTCATAATCACCATCCATGGGATCCGATG
>Gm20_26485526	TTCAAAAACCAAGAAAATCTCACATTCAAGGAACGTGAGTTACGTGGCTCACATGTAACCAAAAAATAATCAAAATTTTTTTTCTTCAAAAAAATAGCTCTCCACCTTGGCTCCTTGCAAAAATAACGCACACACATCACAGTTTCTTTTCGCCACTAGCACTTCATTGAGT[T/C]CATTTCACAGTTCTATTTCGTACCAATGACAACCACCAATACGAGTTTCTCTCTCTAGAACCAACGAAGCGAAGAAGAAGAGAGAAGAGAATCACGGTTTCGCCATGGACGCGGCATCGCAGCTCGTCTCCTGCGGAATCGATCCCTTCCATCGCGCCTCCTCTCCTTCCCCGCGCCAC
>Gm20_26546196	AGTACTTTGGTTATTTAGAGAAGTTGGGCCATTTGTCTGGAATTTGAATTTGAGGCTATATCTTTCATGTAGGAGTGAGTGAGTGGTCCTCCAAGGAGTTGTGCGTTGACAATTTGGGGCTTG TGGCATAAGGAAAAGACACATTGTGATTCTTGCAATAGTTGCTGGAGTTTTCAACT[G/A]TACACCAAAGAATTTGTTGATTGTGTTTCAATATATGTGGAGAAACCATTAACTCAGGATAGTTGCAATCACAGCGAGCAACTTTTATATTCATGATTGCAGTTTAAATTCACCTTTGTTGGTTATAAATTCCTTCCCATGATTTTATTCATGTAATAAACTGAAGAATTTGTTTAC
>Gm20_26658299	AAGAAAAAGATAACTGATGTGGTCACAAAGGGAGAAATGGAGTGAAAAAGATATGGAAAAA AAATTAAGTGATGTTTATAATTACCCAAAATATATCACAATTTAATACAAGTCTAATATGAATAGAAAGGATTACTTGAGCAGTGGTTGACATTATCACTACCATGGAACAAAGCTC[C/T]TGAGATTTCTTATTTGATGACCAAATATAGAGAGTGTGAAGTAACTAATTCTAGCGAGCATCCTTG GATAAAAGAAGATCATCCACTTTCCGGGCATTGCATAGCTCATCAGCACAATCACCCTCTCAAAAACAGCAGCTGCCCCCATTCCAGTACCTGCTCACAATTGACAACAGCCA
>Gm20_26664906	TGAAGTGAGTATATAAATTCAGACAGAGAAAAAGGGAAAATATAACAAACACATTGATCAAATACCTCATTTATTTCAAAAAGATCAATATCATCAAGCTCTAGACCTGCAGCCTTAACAGCAACAGGAATTGCAGCAGCAGGACCAACACCCATTATGGCAGGATCAACACCAACTGCT[A/G]CGAAAGTCCTACATGACAGCCAAAAAATAATTTCTTCAGATTAATAAAGCAGAATTTTTATCTAATCAATTTCACTTTTTAAGTACAATGAAAACATGATTTATGATTTTAGGATAGCTTACAA TTTATTAAGAGAAGGATAAATATAAAAAATGAAATTCCAGTTCCGTTCTGAAAGA
>Gm20_26666320	CATCAACAGAAATGGTGACAGATTTCTCCTCACCAGTTTTTTGGGTCCACAATCTGTTGGGAAGAAAAAGCTCTATTTCAGCACCCAAAGAACCAAGGAGAGATTCAATATACCCTGTAATGGCCAATACCTTAGTGGTAACTGGGACAATTTTCATCTTTAAATTTACCAGAAGCAGTAG[G/C]TGC GGCAGCTCGCCTGTGAGACTCAACCTATCAACATTATTAAGAAGATTCGGTTAAGAGTTGTTAACATATAAAATAGAACAAAATGTTGAATTAATTTAAACATATATCTAACTTACTGCGGCCTGGTCTTGTTCTTCTTGAACCCCAAGCGCTGTGCAACATTTTCGGAGGT
>Gm20_26702577	CATATTCAAAAAATATTTAACTATCCAAACAATGGGTATGTATTAATCTGGAAGACCTATGTAAACCTAGAATCCAAGATCTAAATCCTTTGCTTCCAAAATCAATTTGGAAAGTTTGAACCCA TAAAAGAGAAGGCATAATCAGTGCCAGAAAGAAAAAATGACAAAGAAAAAGAGT[G/T]CCAA GAAATCACATCAACCACACCAATATTTTCAACAAGAATTTTCACTCAAGTGACGCGGGGATGACGGAGCCTCCCATCTAATGCATATATGTGTCCACCATGGAAAAACTTCAATTCATCGGTATGATTCAGGGCGCTTTGGCTTGCTTTGAACAACAGCTTCAATCTTCTCCATCA
>Gm20_26702622	AATCTGGAAGACCTATGTAAACCTAGAATCCAAGATCTAAATCCTTTGCTTCCAAAATCAATTTGGAAAGTTTGAACCCATAAAAAGAGTGAACCCATAAAAAGAGAAGGCATAATCAGTGCCAGAAAGAAAAAATGACAAAGAAAAGAGTCCCAAGAAATCACATCAACCACACCAATAATTTTCAACAAGAATT[T/A]TCACTCAAGTGACGCGGGGATGACGGAGCCTCCCATCTAATGCATATATGTGTCCACCATGGAAA AACTTCAATTCATCGGTATGATTGAGGGCGCTTTGGCTTGCTTTGAACAACAGCTTCAATCTTCTCCATCACAAACAGGAGTCAGTAATGGAATGACATCGATTGCTTTTCATGTTCT
>Gm20_26702784	AATTTTCAACAAGAATTTTCACTCAAGTGACGGGGATGACGGAGCCTCCCATCTAATGCA TATATGTGTCCACCATGGAAAACTTCAATTCATCGGTATGATTGAGGGCGCTTTGGCTTGC TTTGAACAACAGCTTCAATCTTCTCCATCACAAACAGGAGTCAGTAATGGAATGACA[T/G]CG

	ATTGCTTTCATGTTCTCTTGAATCTGAGCAAAAACAGAAAAATAACATAATTACAA TGGCAAGTTCATTCTTGCAAGTTATATCTCAAATTAAGAATGGCATCAATTACTTTCTGTGTT CATTGAACAAGTTGAATCTAAGAAACATGAAACAGAAAAAAAAAAAAAAAAATA
>Gm20_ 2670654 9	TACACCGTTTTCAACACAATCATATCAAGAATCAACAAGTGCGTAAGAGAATGTGGTACTAA CACTCCTCCAGCGAACAAATACCATGTGCCTACACAAAAACACATTACTAGGATTTCTCATA GCAATGATAAAATGTTAATAGCATTTATAGCTAAGCAAAAACCTTGTGCCTGGAC[C/G]AGA GGTTATACTCAGGCTGTTCAACAATAGGACCAACAAGGTCCAATCTCTGTGCCACAGCCCA GGCCTCGGTGATCTGCTGAGCAGACCACTCGCTAGTGCCCCAGTAGAAGGCCAGCCCCCT ATCAATAACATGATTCATGGCCCTCACGGTTTCCTCGATCGGAGTAGAGGTATCCG
>Gm20_ 2670726 4	GTTCTTGATTGCATCTTCTTTGAACTGGATTTGGATTTGGATTTGGATCCGATTAGAGGGG AAGAAGGAGGAGGAGGAGGAAGGCACCTTCGGGAATCGTCGAATCTAGCAGTAGAGAGAG TGTGAGTGAGTTTGAGAACGGAAGAAAGGGTGTGTGCGAGTGACAAAAAGATGGAT[A/ G]GGTGATCGGTGATCCGTGGAATTTTTCTACGTGTAACTGATCGAGATGGGACTTGGG AAGAGCACGTGACATTCAATTAATTAGTACCACATATGTAGTGACATGGATCAAGATATTAT GTAATGGCATAATTCTAAATTTCTTTTTGAGGTAATAAATACCATACGTCTAAGATATT
>Gm20_ 2683150 0	ACAACCACTGACTAGTGAGTGGTGTTAGTTAACGGCGTGAGAGGGACGGCGAAGACCGAA ACAAGAAAAACGATCAGTTAAGTAACGGCGTGAAAGGGTTGGGCGTTTGTGGGTTTAGTGT CTATAAAAAACCGTTATGTTGGAACCTTCTGAGTCCACTCCACCCAATTCTCTCTCTC[A/G]T CTCTTTCTCTCTTCAACATTCTCTCTCACAAAGCAACAACACACACTGAGAGCGAGGG GAATTCAACTTTCTGCAATGGCGTTTCTTCTTCTTCTGCAACAAAATGGGTCGTCTTCTTG CTTCTGGCGCTGCTTATTCGAGAAGAAGCTATGGCTACACCACAGATCTCTGACCT
>Gm20_ 2683152 6	AGTTAACGGCGTGAGAGGGACGGCGAAGACCGAAACAAGAAAAACGATCAGTTAAGTAAC GGCGTGAAAGGGTTGGGCGTTTGTGGGTTTAGTGTCTATAAAAAACCGTTATGTTGGAACC TTCTGAGTCCACTCCACCCAATTCTCTCTCTTCTCTTTCTCTTCAACATTCTCT[C/G]TC ACCAAGCAACAACACACACACTGAGAGCGAGGGGAATTCAACTTTCTGCAATGGCGGTTT CTTCTTCTTCTGCAACAAAATGGGTCGTCTTCTTCTTCTGCTTCTGGCGCTGCTTATTCGAGAAGAA GCTATGGCTACACCACAGATCTCTGACCTCAGGTTGAGACACACGCACAACTCT
>Gm20_ 2683514 8	GAAGTTGGGATTTTTCTCTTGGAACCTGGCAACCCCATGATGATTGCTGGCGTTGCGAC CCCAACTGGCAACGCAATCGGAAGCGTCTTGCAAGATTGTGGCATTGGTTTTGGCCGAAAC GCCATCGGTGGCCGTGATGGAATTTCTATGTGGTGACTGACCCAGGGATGATGACC[G/ C]TGTGAACCCGAAACCCGGCACTCTTCGCCATGCTGTGATCCAGGATAAGCCATTGTGGA TTGTGTTCAAGAGGGACATGGTTATTAGCTGAAACAAGAGCTGATCATGAACAGCTTCAA GACAATTGATGGTAGAGGAGTCAATGTGCACATTGCTAATGGAGCATGCATCACAATTCA
>Gm20_ 2683524 8	GGCATTGGTTTTGGCCGAAACGCCATCGGTGGCCGTGATGGAAAATTCTATGTGGTGACT GACCCAGGGATGATGACCCTGTGAACCCGAAACCCGGCACTCTTCGCCATGCTGTGATC CAGGATAAGCCATTGTGGATTGTGTTCAAGAGGGACATGGTTATTAGCTGAAACAAGA[C/ T]CTGATCATGAACAGCTTCAAGACAATTGATGGTAGAGGAGTCAATGTGCACATTGCTAAT GGAGCATGCATCACAATTCAGTTTGTACCAATGTCATCATCCATGGCTTGACATTCATGA TTGCAACCTACTGGAAATGCTATGGTGAGAAGCTCCCCAACCCACTTTGGTTGGAGG
>Gm20_ 2683525 3	TGGTTTTGGCCGAAACGCCATCGGTGGCCGTGATGGAAAATTCTATGTGGTGACTGACCC CAGGGATGATGACCCTGTGAACCCGAAACCCGGCACTCTTCGCCATGCTGTGATCCAGGA TAAGCCATTGTGGATTGTGTTCAAGAGGGACATGGTTATTAGCTGAAACAAGAGCTGA[T/ C]CATGAACAGCTTCAAGACAATTGATGGTAGAGGAGTCAATGTGCACATTGCTAATGGAG CATGCATCACAATTCAGTTTGTACCAATGTCATCATCCATGGCTTGACATTCATGATTGC AAACCTACTGGAAATGCTATGGTGAGAAGCTCCCCAACCCACTTTGGTTGGAGGACAAT
>Gm20_ 2683553 2	TGGCTTGACATTCATGATTGCAACCTACTGGAAATGCTATGGTGAGAAGCTCCCCAACC CACTTTGGTTGGAGGACAATGGCTGATGGAGATGCCATCTCCATATTTGGCTCAAGCCACA TTTGGGTTGACCACAACCTGCTGTGCACATGTGCTGATGGCCTTGTGGATGCTCC[G/T]G GGCTCAACAGCCATAACTATTTCCAACAACCACTTCACCCACCACAATGAGGTACTACTACT TTTCACTGTAGTGCTTCTGTTTTCTGTGTCAAACCTGCACTTTTCCCTTCTTGTGTTGCA TCATGGCTTAAGTTGTTGTCTTATTTCTTTCAAATTCAGGTGATTCTGCTGGG
>Gm20_ 2683664 8	GGGAGATTTGTTGCTGAATGGTGCCTATTTACACCATCTGGTGCTGGAGCCTCAGCAAGC TATGCAAGAGCCTCTAGCTTAGGAGCAAAATCTTCTCCATGGTTGGTTCCATGACTTCCAA TGCTGGTGCACTAGGTTGCAAAAGAGGCAGTCAGTGCTAGCATATATTATCATCAA[A/G]CA CAACCATGTGAATGAAAGAGGTCATCCATTATTTTCTTTTGTGTTTATTCTCAAGGTTTT ACCATATTATAGTAGAACATCATCATCATATATCCATAATCCATTGCTTCTTCTCCTCAC ACCTCCATCATCATATCCTGCAGTATTCTTGACAAGTGTACATTAAATTACA

>Gm20_26880242	AACCTGCAATTTAATTTAATACAAGATTTTGGACACATACCTAAATTAGGGCTGAGCTCCAGTTAAGGAATTAAGATGCACTTCAACAATGTTAAAGGAACATGACGTTGAAGCGATATAGGAGACGGCGGGACAGCTTGGGATTGATGAAGGGCACGATCTCAGTGAGGGCGCAGAGG[C/A]ATTGTTGGGTTAGTCAGTGAAGGATTGAGAGAGTGAATGAAGGACTGAGTCACTTTAGGTGCAAAAATCAGTAAGCAAACAAAGACGATTTTGGGTTTGAACCGTCTTTGTACAAAACCTACAATTTACATGTTTTCCATCGCGCCTTTATTTACTTATTATACATAGACAATTATATG
>Gm20_26880264	AAGATTTTGGACACATACCTAAATTAGGGCTGAGCTCCAGTTAAGGAATTAAGATGCACTTCAACAATGTTAAAGGAACATGACGTTGAAGCGATATAGGAGACGGCGGGACAGCTTGGGATTGATGAAGGGCACGATCTCAGTGAGGGCGCAGAGGCATTGTTGGGTTAGTCAGTGAA[C/G]GATTCAGAGAGTGAATGAAGGACTGAGTCACTTTAGGTGCAAAAATCAGTAAGCAAACAAAGACGATTTTGGGTTTGAACCGTCTTTGTACAAAACCTACAATTTACATGTTTTCCATCGCGCCTTTATTTACTTATTATACATAGACAATTATATGGAATGTCGTCGTTGTGTTCT
>Gm20_26880300	CCAGTTAAGGAATTAAGATGCACTTCAACAATGTTAAAGGAACATGACGTTGAAGCGATATAGGAGACGGCGGGACAGCTTGGGATTGATGAAGGGCACGATCTCAGTGAGGGCGCAGAGGCATTGTTGGGTTAGTCAGTGAAGGATTGAGAGAGTGAATGAAGGACTGAGTCACTTTA[G/T]GTGCAAAAATCAGTAAGCAAACAAAGACGATTTTGGGTTTGAACCGTCTTTGTACAAAACCTACAATTTACATGTTTTCCATCGCGCCTTTATTTACTTATTATACATAGACAATTATATGGAAATGTCGTCGTTGTGTTCTCTTTATTACAAAATGCCACCGTGAAGTGTCTTAGA
>Gm20_27300839	AAATTTTCTCTTCTTTGATTCAACCTAACCTTCTGAGACCCTAAATATTAAGATTCTCAATCTCGAGACTTCACAAGTCATAACTATTTTCACTTGCAATCGTTTATTACTCTCAACTGCCCTCTTACTTTTTCTTTTATAAGATACCCCTTTTGCATGCATGTCCTCCACCTCAA[G/T]ACGTTTATTCGTTACAGAAATTAAGCTCAGTGTGTGTGAGTGAGAGAGAGAGAGAGATGGGAAAATGGGATACCTTGCTTGTGTTGTTGTGTTAGTCACAGTGCATGCAACTGCAAGTGCACGTGATGTGCCAAAGGGTGGTGGTGGTGAAGAAGAAATCCTAGGTGTGCATGCAAG
>Gm20_27300840	AATTTTCTCTTCTTTGATTCAACCTAACCTTCTGAGACCCTAAATATTAAGATTCTCAATCTCGAGACTTCACAAGTCATAACTATTTTCACTTGCAATCGTTTATTACTCTCAACTGCCCTCTTACTTTTTCTTTTATAAGATACCCCTTTTGCATGCATGTCCTCCACCTCAAC[G/A]CGTTTATTCGTTACAGAAATTAAGCTCAGTGTGTGTGAGTGAGAGAGAGAGAGAGATGGGAAAATGGGATACCTTGCTTGTGTTGTTGTGTTAGTCACAGTGCATGCAACTGCAAGTGCACGTGATGTGCCAAAGGGTGGTGGTGGTGAAGAAGAAATCCTAGGTGTGCATGCAAGT
>Gm20_27300882	AAATATTAAGATTCTCAATTCTCGAGACTTCACAAGTCATAACTATTTTCACTTGCAATCGTTTATTACTCTCAACTGCCCTCTTACTTTTTCTTTTATAAGATACCCCTTTTGCATGCATGTCCTCCACCTCAACACGTTTATTCGTTACAGAAATTAAGCTCAGTGTGTGTGAGTG[A/G]GAGAGAGAGAGAGATGGGAAAATGGGATACCTTGCTTGTGTTGTTTGTGTTAGTCACAGTGCATGCAACTGCAAGTGCACGTGATGTGCCAAAGGGTGGTGGTGGTGAAGAAGAAATCCTAGGTGTGCATGCAAGTGCACCATGCAGGTGTTGGTGACAAGAAGCACTTTTTGT
>Gm20_27300889	AAAGATTCTCAATTCTCGAGACTTCACAAGTCATAACTATTTTCACTTGCAATCGTTTATTACTCTCAACTGCCCTCTTACTTTTTCTTTTATAAGATACCCCTTTTGCATGCATGTCCTCCACCTCAACACGTTTATTCGTTACAGAAATTAAGCTCAGTGTGTGTGAGTGAGAGAGA[C/G]AGAGAGAGATGGGAAAATGGGATACCTTGCTTGTGTTGTTTGTGTTAGTCACAGTGCATGCAACTGCAAGTGCACGTGATGTGCCAAAGGGTGGTGGTGGTGAAGAAGAAATCCTAGGTGTGCATGCAAGTGCACCATGCAGGTGTTGGTGACAAGAAGCACTTTTTGTGTTGGTGGG
>Gm20_27301847	TGCTTTTCAACATTGTAACGATCCTTGATATCATTCTGGGGCTTTGCATGCATGCATGGATATGTTTTGTTAATGTAGCTAGGGCTGCCAATATTAATTGTTGTTTAGGTTAGAGAGTCTTAGCATTGATTATTTCAATTTGATGTTATTTTCACTTCAAGTTGTTATAATTGTCTTT[T/C]TTAATTACTGTTTGTGTTCTGGTCATTACAGGCTGCATGCATCAACTAGGTAAGTGTGTTGTTGTTAGTTTTGACTCCACTTTTTGTTAGTGTTTTCAATGTCATTTGTTGCGTTTTTCTTTCTTTTTATCCAACAGTATACAACCTAAATCTGTGTATGCGTGGAATTTG
>Gm20_27301964	GTCTTAGCATTTGATTATTTCAATTTGATGTTATTTTCACTTCAAGTTGTTATAATTGTCTTCTAATTTACTGTTTGTGTTCTGGTCATTACAGGCTGCATGCATCAACTAGGTAAGTGTGCTGTGTTGTTAGTTTTGACTCCACTTTTTGTTAGTGTGTTTCAATGTCATTTGTTGTTTCTTTTTATCCAACAGTATACAACCTAAATCTGTGTATGCGTGGAATTTGTTGCGCGC
>Gm20_27706455	CGAGCCTATTTGGTACATTCTTTGAATTGTCAGTGTGTACACTAATAAATAAAACACTATTTTATTTGTCCCTCAAGATTTTCAAGATTTTCAAATATGGAATTTTGTATGATTAAATTTCTCCGAATACTACAATTAGTTTTAAACCTTAATTTAAAACAAACAAATTACAATTAGCAAAAAAATTCTATGAAACGCGGAAAAATCTTTTATTTAGCCACGTAAAAGGTTAAACCATGACATACAGTTTGTGTTCCCATTAGCTATATCAACACAAGAAAG[C/G]CACAC

	AAGAAGGAGTTCCTCTCTCAATATCCAACACCAGCAAAGCGTCTCCACCTCCTCTACTAC TGTCACCACTGCCAATGCCAGTTCCACCGCCACCATAGCCGGTACCACTGCCAACGCCAC TTCCTATCCCAACACCACGTCCTTGTCCACCACCACCTGAACGCCACCACTAC
>Gm20_ 2770648 8	TAACAACCTTAATTTAAAAACAAACAAATTACAAATTAGCAAAAAAATTCTATGAAACGCGGAA AATTCTTTTATTTAGCCACGTAAGAGGTTAAACCATGACATACAGTTTGATGGTCCCCATT AGCTATATCAACACAAGAAAGACACACAAGAAGGAGTTCCTCTCTCAATATCCA[T/C]CACC AGCAAAGCGTCCTCCACCTCCTCTACTACTGTCACCACTGCCAATGCCAGTTCCACCGCCA CCATAGCCGGTACCACTGCCAACGCCACTTCCTATCCCAACACCACGTCCTTGTCCACCAC CACCTTGAACGCCACCACTACCACTTCCTGAGCCTATTCCAACGCCTGAGCCAC
>Gm20_ 2787705 9	TCGCCCCCTCCTCTCGTCGAAACCCGACCCCATCTCCAAATTGCGCTCCGGCATCGTACCAT GTAAAGCCTCGCCGAATATCTCCTCTCATTTTCCACTCTGTTTTCTCTGTTAACCTAATTT TCTCGTTTTCTTTTTTTAAATTTCTCGCAGTTCTCTCACTACTTCGGGGATT[A/G]GCTCC TTCTTCGTCCTATGGCTCTCCAGACACTCTCACTAGCACTCCACCAGGTCAAGTTTTTCG CCTTTTTCACTCCTCATTCCACTTCTCCCCCTCACGTTTGGTAATTTACCATTTTGCCTTT TGGGTTTTTTCTCTTGTGTTTTTTAGAAATGGGTTGAGTTAGACCTTAG
>Gm20_ 2787848 2	ATATCATCAAATTTATTCTACAAGCACATGAGTATGTGAACATTTTATTACATGCTAATATAC AGGAAATATAAGCAGTTGTGTGTTGTATGGACTCTTCAGCAGTTGACATTGTCTTCTCTG TTATTGCAGTTCTATGGATGAAAGGGAATCCTTGAAGAATGACACTCCAATTTG[C/T]AGCAA TGCTTGTAAGGATGTGACAAAGGAGAAAACCTTCTATCTTCAACCAGTGTACCTTTTTCTT CTGGTGTAGGTTTGCCATATGCACCTGAAGGGTGGCCTAGTCCTGGTGATGTATGGGGCT GGAAAGTGGCAAAAAGGATGAACAGTTTTGGATATTTTATTGATAGATTCTT
>Gm20_ 2787962 5	AAGAAAGAAGTTAAACTGGAAACAGGAAGCGTCAGTCACCCCACTGAGAAATCTCTTGGA CTATTAGCTCTACAATCATCGGAGACCGGAGAAGATGCCACAGATAAGACTGGCACCACG GGAAAGAAGTTAAACCGGAAAACACGAAGTGTGGTTCGCCACCTAATAAGAAATCTT[G/A] CGGACTAGTTCAGGGTCTGCCCATTTAACACCTGATCAAGTTGCAAAATTTGATGACTATC TTGATACCTTGGAGGACATGCTTGTATGCCTGAAATTGAGACTACCCATCGGATAATCTA ACTCCTACCATTTTGGATGATAATGAAACGATTGAGTGCTGCAAGAAGAAGCTGTC
>Gm20_ 2907004 6	GATTTTTAGAGAGATCGGACTAGTATAAATGAACGTGTTAATGTTTAGCTTCATTACAACTA CATGGCTATGAATATGAAGATGTCATACAGAGTCATGCTAGTTTTGTTTCACTTTTCATGTT CGTCATACCTTCAAAGCTTAGTATGCGTTGCGATCGGGGAGCGCCTCCTGAGAC[A/G]CCT ATTTCTGTATAGGGCGACCTGGTACGCGTCCACCTAAAACATCAGCTTCTAGTTTTGAGG TAGCCCGACCGGTGCCACCATCATTCAATAGGCCTCGAAACAAGGGTACACCTTATGTTCA CTATCCCTAGCAGGAGTACTTCTTTCAGCAGCAGAAACATGCATGGGGTCTCTCA
>Gm20_ 2966627 1	ACTTTTTTGAaaaaaATGTATTTTTTTTCGGGTTTTATATATGTATTTGAAATTGTTGGAGT GTCCCTCATTGGTGAAAAGGAGCACACATTAATAAGAGCAGAGCATAGTAGCATCTTCATT CATCTCTACAGCACAAAAAAGAAAGAAATCCAACCCGTGAAGAAAGCAAAGCAA[G/C]TAAT AATAATAAAAAAACGTTTACCCCTTTTGTCTGCGGTGCGTGGATGCTGGCAAGAAA ACAAGGAAGGGAGGGTTTCATTGCGGATCTGAATTCGCAGTGTTAGGGCTTTGAAGGGTTTT TCCATTTTCTCAATGGCCACGCCCTTTCCCATTCCCGTCACCGCAGCTCAGG
>Gm20_ 2966799 0	ATCAACAATGATATGTGGAGTTGTGGATATTTTTCTCTACCGCTATTTGATTTATATGGAAT TGCTTTACAGTGTCCAACCTACCTGCTGGGTGGAGATATCCAAGCTAGGGACTACGTTGCTA CAAATGAGGTCAAAGAAAAATGGTGTAGTTGATAATTATGGATTTTCAGAGCAAAG[T/C]ATG CAGCGGGCCCCCTGATACTGAGCATATTAGGGAGGATAATACTGTTGAAGAGTCAAATGGTT CACTTCAATCTTCAGTGAATGCAGTGCAAGACCATGTACCTGTTTCTCCTGATGAACCTGCT GGGGAGCCACAAAAGCACACTTATGCTTCCATTGTATGTACTGGCTAGCCTTTT
>Gm20_ 2966800 7	GGAGTTGTGGATATTTTTCTCTACCGCTATTTGATTTATATGGAATTGCTTTACAGTGTCCAA CTACCTGCTGGGTGGAGATATCCAAGCTAGGGACTACGTTGCTACAAATGAGGTCAAAGAA AATGGTGTAGTTGATAATTATGGATTTTCAGAGCAAAGAAATGCAGCGGGCCCTG[C/T]TAC TGAGCATATTAGGGAGGATAATACTGTTGAAGAGTCAAATGGTTCACTTCAATCTTCAGTGA ATGCAGTGCAAGACCATGTACCTGTTTCTCCTGATGAACCTGCTGGGGAGCCACAAAAGCA CACTTATGCTTCCATTGTATGTACTGGCTAGCCTTTTTGTTAATTTCAAATAA
>Gm20_ 2970709 4	ATTGAAACCTGAATCCTGCGGTATTTCAAGTAAAGAAACATTCCTTGCTTTAGATCATACCAA CAAAATTATATTGGCATTGGAGTTAATTCCTTTGATGATGTTTGTGGTTCTCAGCTTTGGA GGAATGGCGAAAGAGGAAGATGGAGAGGGCCAGACAAAGAGAATTGGAGAAAAAT[G/A]G AACAACCTTCTCTCAAGCTTGACTTTATATATTCTTATATCAAGTGAAAGTTGACATGCTG ATTTCAATACATAATGGGACAATTAGTAGTCATTTAGTGACACTGTATAGCATCTTTAAGAT TTCCACAGCTTATGTTATAAAGGAATCTTTCTCGGACAGAATTTGTGATTA

>Gm20_29985714	GGTGGAGTAGCCGGCTTCTGGGGCGCACTGATAGAAGGCCCGAGAATCGGACGCTTCGA CCACGCGGGACGCGCGTTGCCCTCAGAGGCCACAGCGCGTCCTTAGTAGTCTGGGGA CCTTCATGCTTTGGTTTCGGTTGGTACGGATTTAACCCCTGGTTCATTTAATAAAATCCTAGT[C /T]TCCTACGGTAACTCAGGAACCTACTACGGTCAATGGAGCGCGGTTGGGAGAACCGCGG TCACCACTACCCTTGCAAGGTCACTGCTGCGTTGACCACTCTCTTCGGGAAACGGATGAT ATCCGGTCATTGGAACGTGACCGATGTCTGCAACGGGCTGTTAGGCGGCTTCGCCGCCAT A
>Gm20_29986416	AAATTGTTGAGGATTTCTTCCGAGGATGAGCTTGCGGGGATGGATCTTACCCGTCATGGAG GATTTGCTTATGCTTATGAGGATGATGAGTCGCACAAGCATGGGATTACAGCTGAGGAAGGT TGGGCCCCAACGCGTCGTCCACACCCACCACTGATGAATGATTACGATCAGGATTAAT[A/C]C GGCCCCGACAGTATTATCTTCAATTGAAATTACGTGTGACTTAGAAGAAGAAAAAAGATGA TGATGATTTTTGTTGTAATTTATTTATTTGTTTTGGGTTTTTTTTTAATTTGTAGATTTTTT TTTTATGATGGGTAAGTAGGGATTTAATTTGTAATTGTTATTGGCCGTAT
>Gm20_30319057	AAAGAAAGGGAAGCATCGGTGCCAGATTCTGTAGCGAATCCAATAATCAGTCTCCACCA CACTCAGATCTGCGAGTCTTTCTTGGGTTTCTCTGTGTTGTGTGCAAATTCATGTTCAATT TCAAATTTTCTCTGAATTCAAACGGTTTAGTTTAAAGTGTATGAGATCATCAGAATT[G/T]CCC ATCGGAAACTGGAACCCTAACCGAACTTGATCCGAATCTCCAATGTCTTGCTAGCGCTTT CCCTGCAACCTGCAAATGGACCCGACATCCTTCTCCAGACGCGCGAGTGTTCCCTCCCT CACGCGCCCTCGGCGCCCTCTCCGCCTTCCGCCAGACGCGCCGCGCCTTCGCCGCC
>Gm20_30319132	AGTCTTTCTTGGGTTTCTCTGTGTTGTGTGCAAATTCATGTTCAATTTCAAATTTTCTCTGA ATTCAAACGGTTTAGTTTAAAGTGTATGAGATCATCAGAATTTCCCATCGGAAACTGGAACCC TAACCGAACTTGATCCGAATCTCCAATGTCTTGCTAGCGCTTTCCCTGCAACC[T/A]GCAA ATGGACCCGACATCCTTCTCCAGACGCGCGAGTGTTCCCTCCCTCACGCGCCCTCGGCG CCCTCTCCGCTTCCGCCAGACGCGCCGCTTCGCCGCCAACAAAGCATTCGCGGCC CACGACGCTTACGCTGCGGAGTCCATCGCGCAGCATCCCTGGCGCGTCCAGCGGC
>Gm20_30320625	TAAAGATTTGTTATACAGAGTTCGCGTGTAGTAGCCTTGGCAATGGGCTGTTTCATGTGA GGACTGCGGCCTCTGAGGAGCCGATACCGATTATGAAGTATAGTTTGGTTCTAGGTTGAC ACCTTTGCCTTTGAGGGTTCGGCTTACAAAACGGCATACTGGGAGTTTGCTTTCTG[A/G]GA TGATACAATATGCCTCAAATCCTGATTTGCTAGTTCCCTTGATGATGTTACATTCACCTCTG AAACTGCCTGTTGACCCGACACTGTTGAAGGTTTCTCCAAAAGCTGTGTTAAATAGGACTG AAAGAGAAATTAATGGCTTGTGCCAGAGATTCCATTGAAGGGCTCTCCTGGCAG
>Gm20_30451031	AGATAAATGCTTCATACAAATCAGAAAGAAGGGATAGAGAAGCATGATAAACACAAATGTTT GTCACGGAGGTTAAAGGTTTTACAATGTCTATAAACAGTTACTAAAGAGCAGTGAAGCACC ACCTGAAACAATTATTAACAAGGGGCCAAAATGCCTGCTGAACCTCATGAGCTCTA[C/T]TT ATAATCACATTACACCTGAAAAGCTAGGATAAACAAGATGGGATAATACAAGATAGTAGAA CAAAAAAATCTAAAGACAAAAAATTGAATGCCGCTATCTACTTTTCAATTACGGGATTGCAAG TTGAAGCACATACACTTATCTTTTCATGGCTTCACCCCTCACTTCCATACTCTTTA
>Gm20_30452416	ATACATCTCAGAAGCCCTCTGAAACCTACCCAGATCCATATACAACCCACCTAAATTATAAA GTGCATCCACATGACCAGGCTTCAAATCAATAGCCTTCTGAAACACCTCAATAGCCCTCTC GTCCTCGCCCATGGCATGCAAAGCCGAGGCCAAATCACAATGTGCGTCAGCATAAT[A/C]C GGCTTCATGAAAATTGCCTCCTCCAACGCCTTAACAGCTGCCCTATACTCCCCAACACCAA AAAGCGCACTACCTAAGAGCTTCAAAGCTCTAAAATGCGTGGGACAAAGAATCGCGGCCCT CCCTATAATACTCACAGGCACTCAAAACCATCCCTTCGCCCTCAAGTGCGATGCCGAG
>Gm20_30452443	ACCCAGATCCATATACAACCCACCTAAATTATAAAGTGCATCCACATGACCAGGCTTCAAAT CAATAGCCTTCTGAAACACCTCAATAGCCCTCTCGTCCTCGCCCATGGCATGCAAAGCCGA GGCCAAATCACAATGTGCGTCAGCATAATCCGGCTTCATGAAAATTGCCTCCTCCA[T/G]CG CCTTAACAGCTGCCCTATACTCCCCAACACCAAAAAGCGCACTACCTAAGAGCTTCAAAGC TCTAAAATGCGTGGGACAAAGAATCGCGGCCCTCCCTATAATACTCACAGGCACTCAAAACC ATCCCTTCGCCCTCAAGTGCGATGCCGAGGTTGACGTAAATCTGGGGGAGCAAGTA
>Gm20_30453928	TTGATGATAATCATGATAGGCGGTTGTGATGTGGGGTTACTACTCAGAGGGACCATTTCAA AGGGAGCATGATTTTTGAAATTGGAATTTGGAACCCTAATTTGTTGTTTGAATCTTTTG ATGAGATTTGGGGTTCGTTAGGGCAGGGGTTGGACTTGTTTGTGTTGTTGATGATGAT[A/T]AC GATGAGTGGGGCGTGTGTGGGTGAAAGCTAAAGGGGATTGATGGGACCAATTCCAAAGG CTATGATTTTTTTCTTAGGGTTTGAATTTGAAAACCTAAATTGCTTATTTTGAATTCATTA GGAGTTGATTTGGGACTTAGTTTGCACAATTATTAGCTGTCTGTTAATCGTTAT
>Gm20_3052866	CATTGTTTATGCGTCAATTTTCTACACAATAATTCCATACCTTTGCTATTGTACAATTCTCTTT GCTCGAAAGTCAAATAACCATTTAAATAAATAAACTACCAGTGCATATTAATAAGAAATAA

2	GAAATGGCAGATGGAACTTGAATATATTTTTACGTTGGACAAGGTAAACAAT[T/A]TCATCA CTTCCTATTTCGTTATAGTTTGTATCTATTAACGTAATATCCAGAAGCAAAAGGAATTG TTTTTCATCATAAGCTGTCTACTAGAAGAAGTTGATGTCTATCACCCATTTCTTTTAATGTCA ACAAAACACCCCGAACAAAGAATAGCTACAACCATATAAATTTCCGATA
>Gm20_ 3052879 4	GCAGATGGAACTTGAATATATTTTTACGTTGGACAAGGTAAACAATGTCATCATTCTCTAT TCGTTATAGTTTGTATCTATTAACGTAATATCCAGAAGCAAAAGGAATTGTTTTTCAT CATAAGCTGTCTACTAGAAGAAGTTGATGTCTATCACCCATTTCTTTTAATGTC[A/G]ACAAA ACACCCCGAACAAAGAATAGCTACAACCATATAAATTTCCGATAGTAAGTATTGGCCGACAA ATCAGTTATCTCACTAATTAACAATTGACGTTACAAAATTATCTACGGACAACATAAGCTTT GAGGTTTTGATGACTAAGGCAAATATTGCAGCTTTCTATCACTAAGGTTAG
>Gm20_ 3052922 9	TGCCATGTGAGGAGCAAGACTGAATAATCAAAGAAACCACTACCTTCATTAACTTTTCCAAT CTCATCATAGATATTTAGACCAAGCACCAATAGCAGGTAGCAACTGTTTACTCCAGAGCAA GACTTTGGCTTGTAACAAGTGGCATTATGACTCAGTAGGAGCATCTTCACCCGG[A/G]CA AGCCGAATATCCGAAGATTTCCGGTCCATTGATCCACAGCTATGTTTTAGAATCTGGACC AAATTTTACACACGTGTTCTTCCCTGAGAAGAAAAAGTTCGAGTTGTGTATTAGTAAGTAA CAAACATGAAATCATCAAAACCTCAAGGAATAATGAACAATGGGAAAAAGGTATA
>Gm20_ 3053429 4	AATTTAACCCCTATTTCTCTTCGGCTATAAAAAGTCAAAAACAACCTTCTACACAAGTGCTTAC ACAATAAGTTTAAAGTTGTTTATCCAAAAGGACCCCAAATAAAATGTTTCTAACTTCTAAGCA ATTAACCTACAGTAAAGGATAGCACTTTACCTTATCTGCTGAAGCAGTTAGG[T/A]ATGAT TCACCCTGAGCTACAACTTTACACTGGTCACCTAACAAAAGCATTCAATCAGCAGAAGT TGAAGAAAAGATAACTGTAGCCATTTCAACTAAAAATAAAACCTTTTTAGAGTGACCACTGAG TGATAGATAATATCTGTCTGATGGGCGGTCAAAAATAACAGCATTGTATC
>Gm20_ 3053711 3	TTTCTTTCTCCTTTTCAAATTCTCAATTAACACTTTTCGCTACTTGACCTCTACTATTTCTCT CACTTTTTTTGTTATTTTCTCAGTAAATTGAATAAAAAAATGAACAAATAGAGAGAGAAGGG TTACCGAAATGTGTCGCTCAATCAAACGCTTCTCGAAGAGGAGACCGAGTT[G/A]CGAGA CACGACGGGCTCTTCCGGAACCTTCGCGGAGACTGCAAAAACAGAGAAAAACAACGACGT AGAGTCAGATCAGCAAAACCCAACGAATTTAGTATAACAAAAAATCAGAATGAAAGAGAG CGTGTAGTGTAATAATCAGAATCGGAGGAGAAAAGGAAGTAAGTGGAGCAATT
>Gm20_ 3069148 5	TGTGGTGAAGTGTGCGACAACACTGGGGCCAAGAATCTTTACATCATATCGGTGAAAGGAA TCAAGGGTGCCTCAACAGATTGCCATATGCTTGTGTTGGTGAAGTGTGCTGCTACTGT GAAAAAGGGGAAGCCCAATCTTAGGAAGAAGGTATTGCCAGCAGTTATTGTGAGACA[G/T] CGTAACCCGTGGAGAAGAAAGGATGGTGCCTACATGTACTTTGAAGGTAATTCTCTCTCTC AATATGTTAATTTTATTAGAAAAGAACATGCTTGCATTTTAGTGAACTTTACAATTCATACTT CTAGTGTTTTTATTCACAAATGTGTTGTTTCGTACTTTGTTCTGGAAAATGCATT
>Gm20_ 3071399 1	ACTGAATGTGACATCTCTAGAGGAATGGATGAGCAAGGGCAATGCACCTGCTCTTGATCAC TCATTGAAGCTATACAATGAACCTAAATCCAGGGGGTGCAAATCATTTTGGTTACTTCAAG GAAGGAGCATCTCAGATCAGCCACAATTGACAACCTTGTCGAAGTTGGCTATTATG[A/C]TT GGACTAAAATTGTCTTTAGGTGAGTACTGTTTAACCCCTTATCTCATAATTATCTTCTACTAG TTCTCATAATTATCTCATATATATATATATATATATATATATATATATATATATATTAATCTCTGTG CAGAGATCCTGCTGATGAATTGGTGTGAGTGCAGTCAAAAGTACAAGTCTGATG
>Gm20_ 3111466 1	CATGTCAAGGGAGAAGACAGACGTAATTTTGAAGTAGTTGTGAAACACTTCTTTATAGAAA AAGAAGTGACTTCTACTTTGGTAATGGACTCCCTGTATAGTGGGTGAAGGCTCTCGAAGG CCAGAATAAATGCAAGAAAGGTAGGGTAAAATTGTTGGATGCTGAAGAAATGCCCG[A/G]A CCAATTGTCGGTGCTGAGAAAGATATGTTTGTATTGGTGGATGATGTTCTTCTGCTACTTGA GAGAGCAGCTAAAGAACCTCTGCCTCCGAAAGATGAGAAGGGTCTCAAAACCGCACAAA GGTGAGAAAAATAATGTGTTGATAAAAAGTAAACGAGGAATTATGTTGCTAGCACTT
>Gm20_ 3112368 6	CAAGGGGAACCTTTTTCAAATTACAAAGTCCAAAAGTCACTAACAGGTAAGTATAGTG TAGTGTCTTTCTTTAGGTGGATGAGTATCTATGATTAGTCTGAAATGTTAATTTGATTGGCAC TGCTTATTTGTAGAGGTAAGAACCAAGTAAAAGCATCGTGTAATGTGGACAGTT[T/G]CACA ACAGAAATGGATAGTCAGCCGTCCGGTCTTCTGCAGATGCTGTAGATGTTAATGAGTCTG GAAGTAGCAAGTTAGGTGGATCTGAGCCTGAGGGTGTCTGTTCTGCTTGCAGGATCGATT AAAGTGGCTTGATCAGCCTGTCATCAGAAAGGTGGTATTATTTCCAAAAGCTTA
>Gm20_ 3112523 8	TGGTCCGTTAGGTCCTGATCCCTCACCTGCTACTCATAGTTATGTTCCCCAGTCCATAGAA ATGCAATAATGGGGAATCCTGTGGTGTCAACTGCTGCCAGTCTTCCCTATTCCAGCTCCAG TTCAGGAGTAAACCCATCTCCTGGCTATTACATCCACCCATGGTATCATCCCCAC[G/C]AT TTATCTCCCGGAGCTCCGACAAAATGGATTCAAATACTTCCCAGTCTGGTGTTCCTTTGGT ATGATTTACGGGATGTTTTACAGAACGGACCTAATTGGATTGATAGTTCTCAAAGGGAAG



	CCAGCAGAAGCATGCACTATGAACCTCCTTCCCGACTAAATGATGTTCAAAACCT
>Gm20_31543069	CGCCAAGGGGTTGAACTGCATTTCTTCGGTGAATGCTCCTTGGGACCGGTGGGAAGGGC GAGCATCTCGTCACACATAAGGCCCTGGCTGAAACCACCATGATTGAACTCGGAAAGCGT GATGGTGGTTCCCTTCTTCTTCTTGGGCTTGGAGTTCAAAGCCTCCTTGAGGTTGGGAA[C/ T]ACTTTGGGGCTCGACCGCCACGGCCTGGGCCTCAGCCTTCTTGGCCTTGGCCAGTGTG ATGGTGGTTCCCTTAGGTTTTTTGTTTGTGAGAGATGGGAGCATAACATGGGAAAGGTGAG AGAGCTGTGCCGCGACACAATCAATGGGAAGGTTTTTTAAAGACTATTTTTGGAAAAATAAT
>Gm20_31624554	GAAAATGCTCACCACCATATTGAAGATTTCAAATTTTAAATGTGTTCAATTGAGAAAAGAAT CTCCACATTGCAATTGCCACTCCACACAATCTTCAAAATCAAACTGCTTGCACAATCCAGT AGAGCATTTTGTGGTGCTCATACCGGCATATGTAATCAACCCAACAAGAATT[G/C]TGCA GATTGATTCTCAGTCTTTTCCCTGTCAGCAAAGTATAATGTCTGAAGTAGCAGAACATTTG TCCTGCTAACGATTTGGTGTGCTCAAAATTACGCTCACTTTGCCATCTGATCATGTTATGG GCAAGCGGGGCCAGCCACTTGAATATCCCATCAAGATTCTCCTTCCAATCAT
>Gm20_31793087	CCTTGAACCCAAAGGCGAGAAAACGAAAACCCACATCCAAAAATCGAAAACAAAGTGAAA GTGAACTCCTTTTTACGAATCCGATAGATCTGTGAAAAGGTCATTACATCTGCCATGCAGT GAGGATGAAAAGGGAGGCAAGAGAACACGACCCAGAAAGGAGGAGAACCCCATTT[G/T] ACTCCCTTTGGCAGAGGCCAAGAGAGTGAAGCTCGGAGAGAGAATAACGGGGAAGACAGT GAGAGAGAGAGTGAAGCAAGGAACACGAAATGAAATATAATATTGTTGGAGAGAGAGAGG AGAGAGTGGCAGTGACCAGAACAAGGTGGTCAATTTGATTAAATTTAATGTATTAAATTA
>Gm20_31991811	TTGGCCAAGATTGTAGTAAGTTCTTGAAGAGCATTCACTAAGTTGATTGTTGTGTTGTAGTG TGATTTACTAGACTTGCCTGCTGTTACACAACCTGGAGAAGGATGCCAAATATGCTCTGTTAT ATCAGCTTCTTAAGATTTTTCTTACCCAGAGGCTGGATGCATACTTAGACTATCA[C/G]GCTG CCAATTCCACCTTGTTGAAAAGTTATGGTAAGCACTGTTTATGAATGATTATCATTTATCAAT AGTTACATATACAGGGTTCTTTTCTACTTTATGTTACTATCTAAATGTTTTGATGTTATGATG CTCTTTTAGGCCTTGTGCATGAAGAATGCATTGCCAAAATGAGGCTTTTG
>Gm20_31993488	TCCAGGTGGAATTTTTTATCTTAAACAGAGGATAAGAGTGTAATTTGCTTTGTTGCAATTTA TTTTTCCATTATTTTATGTTGGCTGGTGACTCTAATTTGTAATAAATTTGGCAGAGCCATTT TGATTGAAGTTTTTCCATTTTTAATTTAGACGACCTCATTTCTAAATATCTTT[G/C]CTGTGCT CTAAGTTAACAGTGCGATTGTTGTTGCCCTGATTGCATAGTCAATTCAACTATCAAATTC TATTAGTATATTTTCACTCTTACAACGCGTGCTTGACACTTCTTTATACTTGTATTTATAGGA GAAGCGTAAAGTGCAATTTTGAATTCGCTTCAACACTAGACTAGATAG
>Gm20_32068827	TTAATAGTTGAAAGCTTAGCCCTCAGAATCTCCTCCATGTCCTTCCTAGAGTCATTTGCGAA GCTTTTCCTGCTCGTGCAAGGCACCGATCGCCATATTTTCCTGCATGCCCTCCTCATTCTT GCTTCTAATTTCAATCTTACCAATGCGCATATCCTCTAACACATCACACATTAAT[G/C]TAA CTATGCCTATTGATGATACCTTGAGTGACCTTATGACATCTTTGAGTTTTCAAATATAGTGCT AACTCTTCTCTCTATATATATTACTAAAAAAGGAAGAGAGGGTATGTGTATGTATATGATCAA GGAAATTTGATGAGTAGTCAGATTAGTTGGAAAGCTAGCAGCTGTTGAAA
>Gm20_32088857	GGTCATTGCATACAGTGAGAACGCACCCGAAATGCTGACCATGGTGAGCCATGCTGTCCC CAGTGTGGTGACCAACCTGCCCCTTGGTATTGGCATTGACATAAAAACTCTATTCACTGCA CCAAGTGCTTCTGCATTGCAGAAGGCTCTAGGATTTGCAGAGGTTTTGCTTCTTAACC[T/A] TGTCCTTATCCATTGCAAGACCTCTGGGAAGCCCTTTTATGCGATTATCCATCGTGTCACTG GTAGTATGATCATTGACTTTGAGCCAGTCAAGCCGATGAAGTTCCCATGACTGCAGCAGG TGCCCTTGAATCTTACAAGCTTGCTGCCAAAGCAATTACCCGATTGCAATCGTTGCC
>Gm20_32088887	AATGCTGACCATGGTGAGCCATGCTGTCCCCAGTGTTGGTGACCACCCTGCCCTTGGTATT GGCACTGACATAAAAACTCTATTCACTGCACCAAGTGCTTCTGCATTGCAGAAGGCTCTAG GATTTGCAGAGGTTTTGCTTCTTAACCCCTGTCCCTTATCCATTGCAAGACCTCTGGGA[A/G]G CCCTTTTATGCGATTATCCATCGTGTCACTGGTAGTATGATCATTGACTTTGAGCCAGTCAA GCCGTATGAAGTTCCCATGACTGCAGCAGGTGCCCTTGAATCTTACAAGCTTGCTGCCAAA GCAATTACCCGATTGCAATCGTTGCCCTAGTGGGAGCATGGAAAGACTATGTGATAC
>Gm20_32093107	CAGCTTTTATCCTCTCTGCTGAACCTTGACGAGCCCATAAATTTGAAAGGTTAAAGTTTTGG AAAAAAGAATAAAAGATGATGCAACCAGCAAAACACAATCTGTTTTGTACATCAATAAGAGA ACGGAGCTAGAAATTAATTGCATCCAATGTAACCTCTAATTTGTTCCACGTTTCT[G/T]ATTA ACTGCTTCTTGTACGTACGTTGTAGATATGTATTAATCTTCTTCCCATTCGTTGGCCTGTT CTTTCTAGAGTTTTTCTTATTTACTTATTCCTCCATCAAACAGCAATGATGAATCATTTGTT GCGAAATGAAACGTAGAAATTAGAAACCGCTGATTCTGTCTCTCGCTTCTT
>Gm20_3210044	GATGGCCTGTTGTTATTACCCGCAATTCAAATAGATTCTATGTGTGCTATGCTTTTGCAGGT TCTGGAAAGAGATGGGAAAGTCTAAGGAAGTCAAAGAAATATCCAAATCTTGCACGATGGT

9	TCAATTCAATAGTGGAAGAACATGGTACTGCCTTAAATGAAGTCACATCAGTATAT[G/A]CTG GCAAAAAAGGATTGGGTGAACCAACAGCAACCAAGTCAAAAGAACAGTCAGTAAACACTGA TAAAGTGAAGAAGGTGAATGGAGATGTGTCTGAGAACAACAAAGGAGGAAGCAAAACCTTCA GCAGAAATAGATCTTCCAGATGCTGAAGTTGGAAGTTTCGCTTGCGATTGTCAC
>Gm20_ 3210046 7	CCCGCAATTCAAATAGATTCTATGTGTGCTATGCTTTTGCAGGTTCTGGAAAGAGATGGGA AAGTCTAAGGAAGTCAAAGAAATATCCAAATCTTGCACGATGGTTCAATTCATAGTGGAAG AACATGGTACTGCCTTAAATGAAGTCACATCAGTATATGCTGGCAAAAAAGGATTG[G/C]GT GAACCAACAGCAACCAAGTCAAAAGAACAGTCAGTAAACACTGATAAAGTGAAGAAGGTGA ATGGAGATGTGTCTGAGAACAACAAAGGAGGAAGCAAACCTTCAGCAGAAATAGATCTTCC AGATGCTGAAGTTGGAAGTTTCGCTTGCGATTGTCACCTGAACCAAGTGGTTATC
>Gm20_ 3210088 3	CAGGGTCAGGTTATTGTGCGTTTTGATGATACCAATCCTGCTAAAGAAAGCAATGAATTTGT GGACAACCTGATTAAAGATATTGATACATTGGGCATCAAATATGAACAAATTACATATACAT CAGATTACTTCCCTGAGTTGATGGAAATGGCTGAAAAATTAATTCGCGAGGGTAA[T/C]ACA TATGTTGATGACACTCCACGTGAACAAATGCAAAAAGAGAGAATGGATGGCATAGAATCTA AATGCAGAAATAATATAGTAGAGGAGAATCTAAACTGTGGAAGGAAATGATTGCAGGAAC AGAGAGGGGATTGCAGTGTTGTGTCCGTGGCAAGTTGGATATGCAGGACCCAAAC
>Gm20_ 3210096 2	ATATTGATACATTGGGCATCAAATATGAACAAATTACATATACATCAGATTACTTCCCTGAGT TGATGGAAATGGCTGAAAAATTAATTCGCGAGGGTAAAACATATGTTGATGACACTCCACG TGAACAAATGCAAAAAGAGAGAATGGATGGCATAGAATCTAAATGCAGAAATAAT[A/C]TAG TAGAGGAGAATCTAAACTGTGGAAGGAAATGATTGCAGGAACAGAGAGGGGATTGCAGT GTTGTGTCCGTGGCAAGTTGGATATGCAGGACCCAAACAAATCACTTAGAGATCCTGTATA TTATCGTTGCAATCCAATGCCCCATCATAGAATTGGATCCAAGTATAAAGTGTATC
>Gm20_ 3210183 1	CAAAAATCTCAATCTCATGGAATGGGACAAGCTCTGGACCATTAAATAAGAAGATTATTGACC CTGTCTGTCCTAGACACACTGCTGTCATTGCAGACAGACGTGTGTTGTTGACTCTCACCGA TGGTCTTGAGAAACCATTTGTCCGCATCATACCTCGGCACAAGAAATATGAAGCTG[G/A]CG GAGATAAAGTTACAACATATACTAAAAGGATATGGCTTGACCATGCTGATGCAGTGCTTTA TCAGCAGGTGAGGAAGTAACATTGATGGATTGGGGAATGCCATAGTGAAGGAAATAGAG AAGGACCAAGATGGAAATATCACAGGGTTGAGCGGTGCTTTGCATCTTGAAGGATC
>Gm20_ 3210353 9	GACTTTAGCATATGGAGACTCCAACATGCGAAATCTTCAGCGTGGAGATGTATTGCAACTG GAGAGAAAGGGATATTTACAGGTGTGATTTACCCTTCATTGCCCCCTCACAACCAATTGTGC TATATGCAATCCCTGATGGTAGGCAACAGACAAGTTTGAAGTAATCTTTTTATCTTG[A/C]AA CTGTGTGGAGCATAACCCACATTACACTCAGAAAAGTTGGTAATGGGGATAACAAGTTTG TGGAATACTTTTATGTTTACCATCACATTGTACTATGTTTATATGCAAACCATATATCTGG ATTTTTTACTAGCAATTGGAAGTCTTCCCTTTGTTATTTTTGGGACAAATTG
>Gm20_ 3210357 3	CTTCAGCGTGGAGATGTATTGCAACTGGAGAGAAAGGGATATTTACAGGTGTGATTTACCCT TCATTGCCCCCTCACAACCAATTGTGCTATATGCAATCCCTGATGGTAGGCAACAGACAAG TTGAAGTAATCTTTTTATCTTGAAACTGTGTGGAGCATAACCCACATTACACTCA[G/A]AA AAGTTGGTAATGGGGATAACAAGTTTGTGGAATACTTTTATGTTTACCATCACATTGTACT ATGTTTATATGCAAACATTATATCTGGATTTTTTACTAGCAATTGGAAGTCTTCCCTTTG TTATTTTTGGGACAAATTGCAATATAGCCTTAACGAAAATTCTATTGATTTTA
>Gm20_ 3211130 5	ACCGGCCTTGTGATGACTTGTCTTTGCCACTTTAATTTTTTCTATTGCCTAACATTCATATA GAAATCATTTAAACCTTTCTGAAGAGGCAATTGAAATCCTACTCAAGCAGTTCATAATAATA CTATTTAGAGCTCTACATACCTTAGTCACTCTATCACTGTTGAAACCATTC[C/G]TAACC AGAAATGTTATTAACCCCTGTAAGAATGCAACATAATACAATCTAGTTATCTTGCCAAAGAA CACAATATATATAGATAGAACAACATATGTATGTCAGTATGTGAATCCATGTGTATTTGAAA AGATATTAGAAGTTCTCAAAGAGAATAGTTATTACTTCTTCATCTGGAGA
>Gm20_ 3212998 3	AAGAAGGACATGAGAGTTAAGGCTAAAGAAAGCAGCCAAACAAGTGAAGCCAATACTGAG GGACCTCTTCTCTGATGAAACAGATGATGGAAGGAAGTCTAGGGAATGGAAGCACAGC TAAAGTCAGGAGGCTCAAGACATGATTTGGTATTAAGGACTAGATAAGTAAAAACATGT[T/C] JATGTGGAACACTAAAATAGATAGTGGTTGAAATCTCTCTTATCCTACATGTGTCCTTCTTG TATGATTGATTAGCACTGTTGGTTCTGGCTCCATCCGTGTGTCTGTTGTGTGCAGAGTCAAA GAGCTTTTTTTTTTTTTTTTTTCATTTTAACATGTTCTTTTTTGTATATAAGTTTA
>Gm20_ 3214638 7	AGTGTTAAGAAGAACACGTTAAAAAGTGTGAGTATTATCTCGTTTAGTTTATGATCAGACAA TAAATGGTTGGTTGAAGTGCAAAATTAATCAGCCTTTCCATTATTTAGTAAATTGGTGGGA GTAGTTTTCATGTATAGATTACTTAGTTGGGAAAGCTGAGGCAAATGCCTTCGGT[A/C]TTTT TACAAAGATTGGGAGAGGTAAAGGACTATTAGAATATATACACTGTAATGGAGTAAGCATT GACAGTAGTTGAATGAGATAATCTCAATCTCATATTTCCCTTTCAATTTGGGAATGAAAAATA

	TGTCATTATTGAATTATCAATGTTTATTATATACACTGATGACTTAAATAAT
>Gm20_32208099	ACCCCCCACGTGTTGTCAATGAAGTTTGTGCAGCTTTGTAAGTGCATACTCCCATTTGTTTT GTTTGTGCTGTTGCTGTTATTTAAGTTACTTGTTCAGTTTTGAACCCCATCAGATGAAAT GGAAGGGTGTGTTATTAGATTGGAGCTTCACGTGTTGTGTTGGGAAAATTGA[T/A]GGGA GCATGGAGGAGGTTATGGTGGAATTGTTATTAAGGTCTGAAGAAGTGGAGGCTGGAACAA AACAAAGAGAGGCGTTGGGGTTTGCAGAGAGGGACATGTTTGTGTTGTTCCATAGTTCAA GTGCTGGAAGGAGAAGTATGATCCTACAACATCAGAAGTGTGTTGTTGTTGTG
>Gm20_32208274	TTGAAGGGAGCATGGAGGAGGTTATGGTGGAATTGTTATTAAGGTCTGAAGAAGTGGAGG CTGGAACAAAACAAAGAGAGGCGTTGGGGTTTGCAGAGAGGGACATGTTTGTGTTGTTCC ATAGTTCAAGTGCTGGAAGGAGAAGTATGATCCTACAACATCAGAAGTGTGTTGTTG[T/A] TGTGGTGGTGGTGGCAGGTGGTGGGGTACCAAACCTATCTAGGAACAGTTCCTGGGTGG AACTAGTTGTGGAACCTTCAAGAGGGTATCTCTCTGATGATGAATTAGGCATGATTTTTGGT TGTTGCTGGTGATGATCAACCAGACTCACTTCATTGGATCCCACAGACGCTTCTGAG
>Gm20_32208399	TTCAAGTGCTGGAAGGAGAAGTATGATCCTACAACATCAGAAGTGTGTTGTTGTTGTGGT GGTGGTGGCAGGTGGTGGGGTACCAAACCTATCTAGGAACAGTTCCTGGGTGGAACATAG TTGTGGAACCTTCAAGAGGGTATCTCTCTGATGATGAATTAGGCATGATTTTTGGTTG[AT]T GCTGGTGATGATCAACCAGACTCACTTCATTGGATCCACAGACGCTTCTGAGTGCTTTT GTCGGTTGAGGGTGGCATCATGTCCTTTTCGTTCTCAACCTCAGAGAGTGGCTGTGGGTG TTTGGGTCAATGCCTCTTTGCCTCAGCTTCTTCTCAGGCAAGAGTTCATAGGTTT
>Gm20_32285027	ACAGAGGAAGGAGAAGAAGTAGGAAAAAGAACCTGTTGAACGGGAAAAAGCGTTACGAAG TTAGAGATAGGGTTTTGGAGGCTGAAAACGGGAAAAATGGAGGAATAGGGTTTTGATCTGAA AATGCGGAAAAAGGAAAAAGCGCAAGTCCAATTACCGGAACATGTACAGCGAGTGAGTG[G/ A]ATGACGGTCACTTGTTGCTTTCGTCACGTCTCTTATCTCACTGTGATTCTCGAACCTTCC GAGGAATGAAGAGAAAACGCTATCGTTTGGATATGTAGCAGAGGAGGCAAATTTTCCACTT TGGGGTCTTTTCCAAAGAATTTTCAAATTAGGACTGTTTTGTAAGTTTTTCTCCC
>Gm20_32307397	GACAAGGTTGAGGATGCACAAAGTGCAACATCACCCCAACAGATCACAAGGTTGAACCT GCTATTACTATAGAAAAGACTGAAGATTCTGTGCCATCTGTTGCAGTTGAAGACAATGTAAA GGAGGAGAAGGAAGTGGTTCCTGACTCACACAGTATCAGTGACATTGAACCAAGTTCA[G/A] AAGGCACCATCCACAGAACCCGCTGCGGAGGAAACACCACCACTGGAAGTAAAAAGGAA CCAAGCAACACAGATGAGACTAGTCTCTGACAGAATCAGAAGGAGGGGTGCTGAAAGAA GCTCAAGTTTTGGAACATGTGCCAAGTGAAGAGAAGCCTGAGCCTGTGTCTACACCT
>Gm20_32307894	AAGAGAACTAGAATTTGAGGCAGAGGTATTGAAAAAGACTAACAACTCTGAAACAGAGAAA CCAGAGCCTGTGGAAGCTGAAGTTGCTGAAAGCCAGAGAGAAGCAGAGAAAGCAGAGCCT GTGGCTGCTAAAGTTGATGAAAAACAGAGTGAAACAGTGAAAGCAGAGCCTGTGGCCG[G/ A]TAAAGTTGATGAAAAACAAAGTGAGCCAGAAAAACAAGAGAAAGAAGAGAAAGCCTAAGA AAGTTGCTCCTTTAACAGAAGCTGAAAGTGACAGAGCAAGAAGGAAATATAGTTGAGGCTCA CCCTCCCAAAGACTCAGGTATTGAGGATGTGAAAGACACTGGCAACTCGGAGCCAGAAGC
>Gm20_32308143	TTTAACAGAAGCTGAAAGTGACAGAGCAAGAAGGAAATATAGTTGAGGCTCACCCTCCCAA GACTCAGGTATTGAGGATGTGAAAGACACTGGCAACTCGGAGCCAGAAGCAGTTCTCGAA AAAACAGAGAAACCAGAACCTCTGCCAAGTGAAGTGGAGGAAAAGCCAAGAGAAACAA[C/T] JACAAGTAGATGAGGAAGTTGCAGAAGCTAGTAAGGAAATGGAAGTAAAGAAAGAAATTGT GATTGAGACTGCCAAGAGTGAAGGAACATCTGACGACACGATCAAGGAAGAGAAAACTGA GAAAGAGGAGGAAACTAGCCTCACTGTAAATGCCACACAAGTTTCATCAAATGCGGAACC
>Gm20_32308472	ATGCCACACAAGTTTCATCAAATGCGGAACCTCAAGAAAATCCTGTGGAAGTCTTCTCTGAA GTTGTGCTTGAGACTTCCAAGAGTGAAGGAGCTCTGGATGACACGGCCAAGGAAGAGAAA ACTGACAAAGAGGAGGAAACTAGTCTCACTGTAAATGCTGCACAAGTTTCTTCACAA[G/A]T TTCTCCTGAAGCAGCAGAAAAAGATTGTTGAAGAGGTTGATGACAAGAAAGAAATCAAGCATT TCCAATGTGATCGAAGCGGAATCAAACGAAGCCGCTGACATCATAAAAAGTCCAGAGCAAG CCTCCATTGATGAGGAAGCTAAAACAGCTTAAACGAAGAAAGAGAGATTATGTTT
>Gm20_32308794	TAAAGCTGACCTAAACGAAGAAAGAGAGATTTATGTTCTCTGCTGCTGTGGAGGAAAAGGTT GCTAGTTGATGTTGCTGCAATGATGATGTTGACAAGAAGGAACCTGAAGCACCTAATTAATG CTGTCCCTGAATCCTCTCAAGAGGAAGTGGTTGAAAGCAAAAAGGTTGAGGAACAGA[C/T] CGAGGCAAAGGCAGCTACAAGTGAAGCAGCAGAGAGTGTAAAGTTGATAAAGTAGTTGAT GACAAGAAGGAAGAGAAAACTGAGACAAAGGTGGAAGAAATCTCTAGAGCTATTAATGAAC CTGTTAGAGAAACATTGGCATCTAAATTTGAAGAGGAAAGTGTGAACTGGAGCGGA
>Gm20_3230900	CAGAGAGTGTAAGTTGATAAAGTAGTTGATGACAAGAAGGAAGAGAAAACTGAGACAAA GGTGAAGAAATCTCTAGAGCTATTAATGAACCTGTTAGAGAAACATTGGCATCTAAATTTG

3	AAGAGGAAAGTGTTTGAACCTGGAGCGGATAAAGGTAGAGAAAAGAACAACACTGTGGAG[G/A]C TGAGAAAACTGAGGTTCAAGCCACCAAGGAAAATGATGCAACAAAAACATCCAAGGACCTT CCAAAAGAACTCCAGCCAAACCAGCTCAAAGCAATCAAATAACATCATTTCAAAGGTTAA GCAGTCACTAGTGAAAGCAAAGAAAGCTATCACTGGAAAATCTCCATCCTCCAAAA
>Gm20_32309193	TGAGGTTCAAGCCACCAAGGAAAATGATGCAACAAAAACATCCAAGGACCTTCCAAAAGAA ACTCCAGCCAAACCAGCTCAAAAGCAATCAAATAACATCATTTCAAAGGTTAAGCAGTCACT AGTGAAAGCAAAGAAAAGCTATCACTGGAAAATCTCCATCCTCCAAAAACCTTTTCCT[T/A]CG AGCCCAAGGGCGATATTAAAGTCAAATAATGGAAGTGAAGTGAACAAGTTTGAGCATTAAAG TATTTACTTTTCTCTTTCTTTTTCTTTTTCTTTTTCTTTGCTGCTGGCGCATAGCATTATAGCATAT TTGTTGGTTTTTATACATTTGTAATTTTTCTCTTGAATTTGTTGAATGCCAAA
>Gm20_32312983	CTCTTGGAACCTGCACCTCCTTGGCCACCAGCTGTCTTTTTAGCACCGCTGATTCTTCCTC AACATCCTTCACATAAAATGTTTTCTTGTTATGGAGGGAAACTATAAGATAAGTAAGAAAA TTATTTAAACAGATAAAAAACCATAAAATCTAAGACATACTTCACATATGGCACTCAG[T/A]GTCG TCGGTGTCTTTTTGTTTTCGCTTCAAATTTTGATTTGGCTCGTCTGAACCTTGGTCTGCACC AGTGCCAGAGCAAACAGAGGAAGAGGCTACTGCTGGCTCCATACTTTTTTCAGCAGCTGTC TGTCCTTTAGTGCCACTGTGCGCAAGAACTTGGTTGGGAGTTTGATCAATCTT
>Gm20_32313759	CACCATAAGACACATCCTGGTTTGCATTACTCGTATTGTTATTTTCAGTTATATCCGATGCTC TTGATCTAGAAGAAACAAAAGATGTTTGACACTGATGCAATGATGGGGGGAATAACTGACT GGTGCTAGCTTTAGGTCTGGCAGTCTCAACCTGTTGAGCTGAAGAACCCTTGAAA[T/G]CAT TCCCTTGTTCTTCCGAAGTTTTATGAGAATCCCTAAACGCCATATTGCCATTACTTTTTTTAT CCAATAGAGTAAAACTGTTTGAGGCATGTAGGAAATCAGAACCATATTCATGTTGCAAGGA ACCATCCATTGATGTATAATCCAACCAAGGTAACACATCTTCATCATGACAGA
>Gm20_32332029	AATAATTGTGATTAAAGGAATTTCAAGTGGGTTTTGGCTTTGATTAGCTTTACTTTTTATGGTG GACCTCCTAATTTCACTGATGAAATGATGCTTTTTTTTTTGTGGCAGGTGGACAATAAAACAA ATGTCATGGGCGTTTTCTGTCTTTCAACTCAGATATTAGTGTTCTTTAATATATC[A/G]JCAAT GTTGTCATATTCTTATGTTCTGTTTTTAAACCCCCTTTGTTCTTAGCTTAAGCGAGAGTGGG GTGTGTAGCTTACTAGCTTGTGTATGTAGTCTCTAGATTTGGTAATTGATTTGAAATACACA ATTTCAITTTGTATCAACGATCTTGTTTCATGTATCTTATGTAAACTGTGCT
>Gm20_32333734	CTTGACATCAGCATACACTATCGAATCATATTAGAAAACAAAGGAAAAGGACATTATCATT GTCATTAATGGAAAAAATAAACTTGCACCTTAACAGAAACTTTGATTTGATCTCCTTACAAGT TACAATGAAGATACAGATTTAACATGAAGGAAACAATACATCTTCGAATTAACC[T/A]TACCA ATTTACATTGCTCTCTGGTCACACTCGAAAAAGAAAAATGAAACCTGAGCTCTCATCCTTTA AACTGCCACTCCTCATCGGCATATGGGGCAATGCGCTTGTGATGTCTGAGAATTCACCCATT TCAGGATACAGTGCAGATGAAATGCGTGATTGCATACACCCCCAAACTGTAGA
>Gm20_32416479	TTATATGTATTTTTGGGCCAAAGTTAACAACCTTACGTTATTGAAGATTGGAGAGGCCAAAAGA AGAAAAAATCCGGGCAGCATTTTATAATCTTGCCAGAGAATGAAAATAAAGCAGAAATTTT AGGTCATCCATTTCTATCCACAATGACATGGCATATAACATAAGGGTATAATATA[G/T]AATC TACTGTTCTGAATGACATTTAATGAAAATGGTCCCTTCAAATGCAAGCACACAATACATCAA GGTGCATAGTTTTACAAGTACATTACAGGATCTATAATGTAAACCTGTGCCATAGTAGTTA CAGAAAGCTTATTTACAATACAATAACGAGCTTATTAACCATGCCCCTATAT
>Gm20_32416493	GGGCCAAAGTTAACAACCTTACGTTATTGAAGATTGGAGAGGCCAAAAGAAGAAAAAATCCGG GCAGCATTTTATAATCTTGCCAGAGAATGAAAATAAAGCAGAAATTTTAGGTCATCCATTT CTATCCACAATGACATGGCATATAACATAAGGGTATAATATAATCTACTGTTCT[T/A]AAT GACATTTAATGAAAATGGTCCCTTCAAATGCAAGCACACAATACATCAAGGTGCATAGTTTT ACAAGTACATTCACAGGATCTATAATGTAAACCTGTGCCATAGTAGTTACAGAAAGCTTATT TACAATACAATAACGAGCTTATTAACCATGCCCCTATATAATCTCCAACATGT
>Gm20_32416571	TGGCCAGAGAATGAAAATAAAGCAGAAATTTTAGGTCATCCATTTCTATCCACAATGACATG GCATATAACATAAGGGTATAATATATAATCTACTGTTCTGAATGACATTTAATGAAAATGGTC CCTTCAAATGCAAGCACACAATACATCAAGGTGCATAGTTTCACAAGTACATTC[T/G]CAGG ATCTATAATGTAAACCTGTGCCATAGTAGTTACAGAAAGCTTATTTACAATACAATAACGAG CTTATTTAAACATGCCCTATATAATCTCCAACATGTCTTGTTAGGCAGCTTTGGCAAAACCA ACTCGGAGATTGCCATAGTCAAAGACGGTGTGATATACCCTCATGAAAACAT
>Gm20_32416629	CATGGCATATAACATAAGGGTATAATATATAATCTACTGTTCTGAATGACATTTAATGAAAAT GGTCCCTTCAAATGCAAGCACACAATACATCAAGGTGCATAGTTTCACAAGTACATTCCACA GGATCTATAATGTAAACCTGTGCCATAGTAGTTACAGAAAGCTTATTTACAATAC[C/T]ATAA CGAGCTTATTAACCATGCCCTATATAATCTCAACATGTCTTGTTAGGCAGCTTTGGCAAA ACCAACTCGGAGATTGCCATAGTCAAAGACGGTGTGATATACCCTCATGAAAACATCCCCA

	AGAATCCTTCAGGCAGATAGTGACAGCAGAATCAGAAAAATAAATGGAGGCCA
>Gm20_3241901 1	TATTCATGAGATTTTCCACAATAAAATATACATTTCCAGAGAAATCCTGATCAAACCTAAGTTT TGTTTTGCTAACCTGCCAGTAACCCTTTTAGTAACTGGAACATAGATATGTTCTCCTTTGAA GTGTTTAGGATCAACACCTCCAAAACTAGTTCACCACCATTTTTCACTTTTG[G/C]ATCACC ATTGAGCCAAAAAGAGAACACCTGTTTCGCTTACAAGATTTTGCTTCACCATATTGTACCTGT AGTTTATAATGTAGCAGAAATAAAAGGACATAAAAGACTTATCATTTTGCTTCTGACATGG GAATCAGATGTAAATGAATTAACAAAGGACAGGATGAAACATACCATACT
>Gm20_3243654 6	CAGTCCCTAAATAAAATTCAATAAAATGTAATTAATGATATTAATAAAATCAACGTAAATT ATACAAATGTCGTCCCCCTGTTATAATACTAACCCGGCCTGGGTCACTTGACTCCGGCACA AGGAAAGCTGCTTCTTCTTTGCAAAGGCGGCGAGGCACCAACGCTCTCTAGCTCC[C/A]GT CTCTTTGTATTTTGAGGTAATTCTTTTACCATTCTGAACTGAAGCTTCAATTCAGCTTCAGCT TCTTCCCTTTCATGCTTATAACGAGTTTCTTTCTGTTAAACCTCATGTTGCTGCAACCTGAGT GCAATTTATTCGTTTAGCCCTTATCTAACATGCTCCTTTAATTCTGTTTCTG
>Gm20_3244004 1	CTTATGTATTCTCTCAATCTTTGTTGGTTATGCAGGTCCTCGAATTACAAATCCTAGGAGTC TTCGTTGGGCAGCATGTGGAGCTGTAAGTGCAGCTCAGGCACGGCTTTGCTGGTTCGTT TGTTTAGTCTGAATGTGAGCCCCAGAATATAGCTGCTTATGACAATAAAAACTAAC[T/C]G GTGTTGTCACCCCCCTTTCTTTTCCCCTACTGATAACATCATAATATACTTGATGAGGTAGG TTGCAATTATCTCGCATGTGTAAGTTTTATGGTATGGATGACATTTTCTGGCACATCACTAA GGGAAAAATTTGGCTAATTAATCCAACCTATTCTTAGATGCTATAAAGGGTAAAAAA
>Gm20_3244014 7	GCTTTGCTGGTTCGTTTGTTTAGTCTGAATGTGAGCCCCAGAATATAGCTGCTTATGACAA TAAAACTAACTGGTGTGTCACCCCCCTTTCTTTTCCCCTACTGATAACATCATAATACT TGATGAGGTAGGTTGCAATTATCTCGCATGTGTAAGTTTTATGGTATGGATGAC[G/A]TTTT TGGCACATCACTAAGGGAAAAATTTGGCTAATTAATCCAACCTATTCTTAGATGCTATAAAGGG TAAAAATTTATTTTCATTCTCTTATTTCTCCCCTGAAAATTTTCTCCAACCATTTTTTCATTCC AAATTAGTCGTTCAACTTATTTATTTACATACCGATTATATCAAGGCT
>Gm20_3244785 3	CAGGAGTCAAAATCCTATGAATAGTTGTTAGTGGGGTGTGTGACAGAAACCATTGAATTTTG CAAGCATAAAAACGAAAAGATGGCAAATGGATTGCAATGATACCTGCACCAACTCTACAA ATATGAATAATAATAAGAAGAAGAAAACAGCAGCATCCATAAAAAGGGGTACTTTT[T/C]TG CTGGTTGGTTCCTTTCCCGTTTTATATTCTAATTATTTCCCATGTCACATGTTGTGCAAT GAAAGATAATTAATATCACACACTTTGTCCACATTCTCATTCTCCTCACTCCCATCATCTTT TCCAATGGCCTCCAACGGCGACGCCGTTTCATCCCCACCCTTCCAAAAAGATC
>Gm20_3244787 4	AGTTGTTAGTGGGGTGTGTGACAGAAACCATTGAATTTTGCAAGCATAAAAACGAAAAGAT GGCAAATGGATTGCAATGATACCCTGCACCAACTCTACAAATATGAATAATAATAAAGAA GAAGAAAACAGCAGCATCCATAAAAGGGGTACTTTATGCTGGTTGGTCTTTCCCC[A/C]TT TTTATATTCTAATTATTTCCCATGTCACTATGTTGTGCAATGAAAGATAATTAATATCACCA CACTTTGTCCACATTCTCATTCTCCTCACTCCCATCATCTTTTCCAATGGCCTCCAACGGCG ACGCCGTTTCATCCCCACCCTTCCAAAAAGATCAAACCTCTTCAGCAATTCATT
>Gm20_3244959 9	ATGCTCTGCTCTTAGTTTTAATTAACCTCTCAATAACATATTCGTCCATTTTATGTGGGAT ACTATTTGCAAATGTTGAGTTTGCACTTTGCTGATTGCGCTATTTTGCCCGCTGTTGAAA ATTTCAATTGTCTATATATCCAAGTATTTGAGTGGACATTTGTTGTTTAACATTT[C/G]CACTAC TCGTTTCTGCAATATTGCACCATCAGTACTGTAGTGGTCTGGTGGTTGCTTCTTTGATATTT TACAACATTGCTGCTGAAGCACAAATAAACTCATGATATCCATAGGTTTACACTATTTTG GGTGATAATTGGAATTTGCAAACCATTCATCTCATTAGATGCTGTGT
>Gm20_3249452 8	GGGACTATTTTGATGAATAACAATCCCCCATTTACGAAGATCAATTTCCAGAACAACCTCTG ACCAATTTACATGAACCTACAACTCCATACAATGAAAGGAACTAATACTAGTTTTCAAAAAATT ATGTACATGATCAATATTAATCATAGTAGGTTGTACAACCTACATACCATAAGC[G/C]GCAAT CATTTCCCTCACCTTCACCTCAACTGAATTTAGCTCATCCATGATCATTTTGGCTGCTCATA ATCACTTAAGTACCTTTGTTCCGTTCTTACTCTAATAACTTTCTGCTCCTACAGACCTACGCA CTGCATCTAATGTAATGCTACATATCATTACAACCTCAACGTATAGCTA
>Gm20_3249520 2	CCATGAAAAAATTGCTGTCTCTCACTTACCCGAAAGGGCCTGTTCCCTGAAGATGAAGATGT TGGTGGTGTGTTGGAATAGATGGTGCAGGTGTTAATGGACTTGGTGGTGGTAGTGTTGTT GGACTTGATTGCGTAGGTGGTGGTGGGGGTGATGATGACTTTGGCGTTGTTGTTGTT[C/A]T TGATGATGATGATGGGAAGATGCACGAACCCGTGCCTATTCACAATTCAGAACAACCAAGTT AAAAATGGGCACTAGTTTAGATTAATAATAAACTTAATTTAAACTTACTGAATAAGTACTTA TTATTTAAGCACTTATATATATATATCTTGTCTTACAATTCAAGTAAAAATTG
>Gm20_3250703	CTAAGCTAACTAGCAAGTCAAATGAAAGTATGAAGGGTGAAATCACTTTCTGATATTTCTTA GTAATGTTTCATAATCCATTCCATGAGGACTCACTTTACAGATACATCACAAATCACAAATCACA

5	ATTGCATGCTTCTCTTCTCTTTTACTGCACTGGAAATTTCCAATTGGATGTATGC[G/T]TATTCCTATTGCCCTTGTA AAAACATCTAATTCTATTCCTTCCACCCACACTCTTAACCTCCACCCTTCTTCTTATCATCCCCACCCAACCTTACATTTACGCTCACATTTCTAAACCCACTAAGGATTTTCAGCTTTTCATCTATTTTTGACAACTGGGTTCTTTGTCTTCCATGTTCTTT
>Gm20_32546952	GGACGGAATGGATTTGCTTTTTTCTGTCTCTTAAATTCTTGTTTAAACGGTGGTGCGACTTTCCTATTTAGTTCTGTTTTTATGAAGCCTGTTCACTGCAATTTTGCTTCTGAAAAAAAAGGATATATATATATATATATTAATTATGGTGGAATTGTTCACTTCTGGCAGGTAATG[A/G]ATGGTTTTTGAGCAACAGCAGTTAAAAGAGAAAAAGGATTACAGCGAAGATGACATCGGTTGGTGTG GCACCAAGCTTCGGGTTTGAGAGAAGCCAGTGGGCATGGAGCAGCAGCTGCGGATAGATTGCCAGAGGAGATGAACGATATGAAAATTAGGGATGATAGAGATGTAATTTGTAA
>Gm20_32551636	AAAATAAGAAATAAATAAATAAATAAATAAATAAATAAATTATTATAATAATAAAAAAAAACACAAATAAATGACTAAGATATCCGGTGGCGACGCTAGATATATCGAGAAGTTTTCTGAAATTTTCAGTTTGTGTTAGTTTAGATTGAAAGCGAGAATGGCAGTGAAGAAGTGTT[C/T]TTGC GAGAAGCAATTCCAAGACACAGCAGCGGATCGGAAACGGCAGCTTCTGGGTATTCAACAGCAGCAGGCCAAAGCTCGCTGGTACGATTCTTCAAACAGCAACAAATCCCTCCC GTTCCCAATCGATCCCTTTGCTTTCACTTCGTCAACACGGTATCATCTCAAGCATTCTGCTC
>Gm20_32551661	AATAAACTGACAATTTTATTATAATATAAAAAAAAACACAAATAAATGACTAAGATATCCGGTGGGCGACGCTAGATATATCGAGAAGTTTTCTGAAATTTTCAGTTTGTGTTAGTTTAGATTGAAAGCGAGAATGGCAGTGAAGAAGTGTTATTGCGAGAAGCAATCCAAGACAC[A/C]GCA GCGGATCGGAAACGGCACGTTCTGGGTATTCAACAGCAGCAGGCCAAAGCTCGCTGGTACGATTCTTCAAACAGCAACAAATCCCTCCC GTTCCCAATCGATCCCTTTGCTTTCACTTCGTCAACACGGTATCATCTCAAGCATTCTGCTC
>Gm20_32551686	TATAAAAAAAAACACAAATAAATGACTAAGATATCCGGTGGGCGACGCTAGATATATCGAGAAGTTTTCTGAAATTTTCAGTTTGTGTTAGTTTAGATTGAAAGCGAGAATGGCAGTGAAGAAGTGTTATTGCGAGAAGCAATCCAAGACACAGCAGCGGATCGGAAACGGCACGTT[T/A]T GGGTATTCAACAGCAGCAGGCCAAAGCTCGCTGGTACGATTCTTCAAACAGCAACAAATCCTCCC GTTCCCAATCGATCCCTTTGCTTTCACTTCGTCAACACGGTATCATCTCAAGCATTCTGCTCTTTTCAATCCTCATTGACTCTCTCTATGTTTAGGGATTTTGCCGTTACGG
>Gm20_32554848	TAGTTTGATCCTAATAAATTAGAATTTAGTGTTCTTTTAGATTCTTAGTTTAATGAAGTTAATT TTAACGAATTCATTAGTTTtagTTTGTTTGTAGAGTTTTGGTACATCAACAACAAAGGAAG AATTTGAGTTGATGAAGCAGCCCCCTTGCACATGCTTAATACGTTAGAGTTA[C/T]TTTGGAAGGTACATGGTAAGTATCTAGAGTTGCTGGGAGAAATTTGAAGTTTGGTGTGTGCTGAGC TAATTTCTCTACTACCATTGATTTGGTTAAGTGCAACCTCAAGCTAGCGCACAGTCAC TTAATCAAGTGTGTTTTTGTATGCGCGTTTAGCGCAGAAGTTGGGCTTA
>Gm20_32559826	TTAAGCTCAACACACTGTCACAGCAAAACCTACCCAAGAACATCCCCAAACAGAGGAAATAGTTAGTGATCTGAATGATAGA ACTCTACTAATAGGAAGTGTAAATGGCTTCATCAGTCCATTGAGCATCTAGCCGAAAAACCACTTTGTGAATTTTTAGTTTTCAATTTTACTCGGACA[T/C]GCG TTTTCGTATACAATTCATT CATAACCAAGTTACAACCAAAAAAAGAACAGTATCTTAAGGACAAGGAAAATGAACATTTATT CACAAGGTAAGACTGAAGATGTCAACATGTCTTGACAATTTGTTGAAGCCTTTAATCAGAAACATTTACTGATGAAAGCAAACGCTTCTGCTACA
>Gm20_32619522	ATAATATATGACTAATACTATTGTATAAGATTAATTGATTGAGTTAATTAAGGAAGTACCTGTTCAGCAAAGGAAGTGGGCTTAGTGACTTGGACAGTTTGAAGGAGGTCACGGTAGGTATCCAGATTGGGCTCATAAACAACATGCCAGCATTGAAATAGAGAGGAGGCTTGGGCCC[T/A]A AGTGAGTGGGCCACTGAACCTTATGAGGGCACTGCTGGCAGTATCCGATTGATATGATACTGCA GAGTGTGGCCCCAAGTGGGCTCACAAAAACAGTCCATCACCGCATAAAAGTAGTTATCAG GCAAGTCAAACAAGTGGTCAATATTGTCAAACACTTGTATATCACCGTCTAGGTATATC
>Gm20_32619566	TAATTAAGGAAGTACCTGTT CAGCAAAGGAAGTGGGCTTAGTGACTTGGACAGTTTGAAGGAGGTCACGGTAGGTATCCAGATTGGGCTCATAAACAACATGCCAGCATTGAAATAGAGAGGAGGCTTGGGCCCCAAAGT GAGTGGGCCACTGAACCTTATGAGGGCACTGCTGGCAGT[C/T]TCCGATTTGATACTGCAGAGTGTGGCCCCAAGTGGGCTCACAAAAACAGTCCATCACCGCA TAAAAGTAGTTATCAGGCAAGTCAAACAAGTGGTCAATATTGTCAAACACTTGTATATCACC GTCTAGGTATATCATCTTGCTGAACTCCACAACTGAGATATACATTTTGCAATCAC
>Gm20_32686866	AAATTTTTTGGCTTACCCTCTTCGAATTTGCCCCCTTACCAATGCATTTTCAGTTTTAGCACA TTTAACAGCGACGTTTCCACCTTCAAGGTTGCAAAATTTCTCGAACTCTGTTGAGACCCGCAT CAATGACAACAATTTTGAGAGGATTTACGTACAGGGTGAATGAACAATGTGAATTTGAA[T/C]CCGT TGGTGGTGAAGGTGTTTCATAAGGATGATGAGAGTGTGTCAGGAGAGAAGAATTTAGGTG GTGATGTGAATGCGAGCGTGGGAAAGAGTAAAGGGGAGGACTCAGAGGTTGGAAAAAGAA

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>Gm20_ 3268995 4	ATGTATCAAACATGGACCATAGCTGGTTTCCTTACATCCAAGATGCTTTTGAAGAATCCTGA GATGGCATCCATGTTATTCTGGGAGGAGGATTACGAGCTTCTTGATATATGTGTTTGTGGA CTTAGCAAGAGTGGCAGGAAAAGGTGTTCCCGTGGTGCCGCAAGGTCTCAAATTCG[C/T]G TATGATGAATTCCTAGGGAAGGGTACATTTCATAGCCAGGGGATCATATATTTATATTGTGCG TTTGGTAGGTTATAGTAAAAGTTAAATTGAAAAATTAACACCTTCTCCTTTGTTCTAGGAGG GCGATTCTCACATTTTTACTTTTTTTTTGTCATTGAGGGGATAGAGATTTAAAGT
>Gm20_ 3269141 1	GAAAAGAAAGTACTAAAAATAAGGGGGGAAAGTAAACCCATAAACAGCTGACAGGTGTCTT GGTTCCATTTACTACCAGTGAGTGGAGTTTGTAGACAGTAGTAGAAAGGGAAAGAGGGTTG ATAGAAGAGAAGGGTTTAGGATTTTGCACAACCTACTCATGGAGGAGGAAGAAGCCC[T/C] AGACGTTATATGCGAGCTGGAACCGTTCAAGGCTTAGTGACGCCCTCACCGCCGTTCTG GTGGAAGCGCCAGCAGGTAAGTAAATAATAATGCCCTAACCCCTAACCCCTAGCCCT AACCCCTAGCCCTTAATTTAATTAACGACCTACCTGGCTAGGACGCCGTTTGGAGTTGT
>Gm20_ 3271063 6	CCTCTTGGCGGACGACAAAACCACGAAGAAGTTGAGAGAAGTCTCGGTCTCCGCCACCT CCGCCCTCTTCCCCGACGGAGACTGACCGCAATTCAAATTCGGCGACGAGGTTGATGC GTTCCACAACGACGGTTGGTGGGAGGCCACATAACCCAGGAATTGGAGAACGAAAGAT[ C/T]TGCGGTGTATTTTCAGAGTGTCCAAGGAGCAATTGGTCTTTTCCAAGGAACAACCTCAGG CTCCACCGGGAGTGTTAAATCACGATTGGGTCCCACTGCAGCAGAAACAACAAAGG CAACAGGTGCCTACGATTCCGCCCTGTTTTCTTCTCATTGCTGCATTATAGAGATTTAGGG
>Gm20_ 3271074 2	GACGAGGTTGATGCGTTCCACAACGACGTTGGTGGGAGGGCCACATAACCCAGGAATTG GAGAACGAAAGATTTGCGGTGTATTTTCAGAGTGTCCAAGGAGCAATTGGTCTTTTCCAAGG AACAACCTCAGGCTCCACCGGGAGTGTTAAATCACGATTGGGTCCCACTGCAGCA[A/ G]AACAACAAGGCAACAGGTGCCTACGATTCCGCCCTGTTTTCTTCTCATTGCTGCATT ATAGAGATTTAGGGCTAACTTGTATATATGTTAATTTGATTCTAGTTTTAATGTAAAAATTTA TTCTTATTTGCTTCGTTAGAGATGACACTTGACTCTAGGGCTAGGGTATTGCTAAA
>Gm20_ 3271103 8	AAATTTTATTCTTATTTGCTTCGTTAGAGATGACACTTGACTCTAGGGCTAGGGTATTGCTA AATTGCATGCATGTTAATTTGACTCTAGTTTAATGTAATGGTGGTTGTGTTAATTACAGGGC AATGGTGAATCGAAGAAAGTGTTGTTGACACCGAATGTGAAATCTGTTGAACTG[G/A]AAA GGGGAAGGGGATAGGTGTGGGGGCGATAGTTGAGGTTAGCAGTGATGAAGATGGGTTCA GTGGAGCTTGGTTTGCTGCCACTGTTGTTGAGGCACTGGGGAAGGATAAGTTCCTTGTTGA GTACCATGATCTGTTGGCTGATGATGATTCCCAGCTTAGAGAGGAGATTGATGCTCT
>Gm20_ 3271104 9	TTATTTGCTTCGTTAGAGATGACACTTGACTCTAGGGCTAGGGTATTGCTAAATTGCATGCA TGTTAATTTGACTCTAGTTTAATGTAATGGTGGTTGTGTTAATTACAGGGCAATGGTGAATC GAAGAAAGTGTTGTTGACACCGAATGTGAAATCTGTTGAACTGTAAAGGGGAAG[T/C]GGA TAGGTGTGGGGGCGATAGTTGAGGTTAGCAGTGATGAAGATGGGTTCACTGGAGCTTGGT TTGCTGCCACTGTTGTTGAGGCACTGGGGAAGGATAAGTTCCTTGTTGAGTACCATGATCT GTTGGCTGATGATGATTCCCAGCTTAGAGAGGAGATTGATGCTCTGCACATTAGGC
>Gm20_ 3271133 5	TGTTGAGTACCATGATCTGTTGGCTGATGATGATTCCCAGCTTAGAGAGGAGATTGATGCT CTGCACATTAGGCCTCATCCGTGGACACGGATGTTGATGGTCAATTCGATTTCTGATG AAGTTGATGCTTTCTACAATGATGGTTGGTGGGTTGGTGTGATTTCCAAGGCTCTTG[T/A]A GATTCAAGGTATGTAGTTTACTTCAGGTCTTCTAATGAAGAACTGGAGTTTGAGAACTCCCA GTTGAGGCTGCACCAGGACTGGATTGGTGGCAAATGGGTGATGCCTTGTAAGGTATGGAT ATTCAGTCTACTCTTTTTACTGTTAAAGTATGTATGTTGGAACCTTTTTAAATTT
>Gm20_ 3273392 7	TAAAATTTGACAAAACACATGTGAAAGCAAGGTGTGGAGCGATATGGATCACAGAACTTAC CATCTCCCCTAGGCACTTGAGGCAACACATAGTATCTGAAGAAAAATTCTGGAATTCGGC ACCTGGCAATGATGATTTTGATTACCATTTTCAATTAATTGTGTTGACCTAACGAC[T/C]TT GCCCTGTTTCTGACAAAGCTCACAAACGTACTGTGTAGATTCACTCTGGTAGACCACAATAA TATCAGAAGCAAGAAAAAGAGAACAAGAACCCAGCAAAACAAATAGAAAAAGATTAGAGT CAAGAAATTTAAGAAGATGGAGAAGGGATAAATTTCTGGAGTACAAATTCAGAA
>Gm20_ 3273395 4	CAAGGTGTGGAGCGATATGGATCAGAACTTACCATCTCCCCTAGGCACTTGAGGCAAC ACATAGTATCTGAAGAAAAATTCTGGAATTCGGCACCTGGCAATGATGATTTTGATTACCA TTTTCAATTAATTGTGTTGACCTAACGACCTGCCCTGTTTCTGACAAAGCTCACA[C/T]ACG TACTGTGTAGATTCACTCTGGTAGACCACAATAATATCAGAAGCAAGAAAAGAGAACAAGA ACCCAGCAAAACAAATAGAAAAAGATTAGAGTCAAGAAATTTAAGAAGATGGAGAAGGGA TAAATTTTCTGGAGTACAAATTCAGAAATGTAGTTAATCCAAGTTATAGGACCC
>Gm20_ 3273618	CAGAGAGATGCTTTCCGGCATTTGAGATATCCGGTGACTTTAACAGCGGATGAGAAGTGA AGTGACAAGTTCATCGACGATCCGAATAGATAGAGAAGTGAATCCACATTCCTGGGCTACT

0	TTGTCGTTTTAATGGCTGATGACAATTGGCGAGCTTGAAATCAGTGTCGTGAAACAGA[C/T]A GTTGTAGGCTTTGAGGGCTTCGATGTGGATCTCGAGAGAGGATAAGTTGCCGCCAACCTG TTAAAGAGAAAATGAGTAGGGAATGAAAAAAGGAGAGGGGAATAGTATAGTTACATACGGT GACATTGACGCCGCGGGAAGAAAGCCGAACACGACCGAGGAGGCTGAGGGAGGAGCAG
>Gm20_32736338	AAATCAGTGTCGTGAAACAGAGAGTTGTAGGCTTTGAGGGCTTCGATGTGGATCTCGAGAG AGGATAAGTTGCCGCCAACCTGTTAAAGAGAAAATGAGTAGGGAATGAAAAAAGGAGAGG GGAATAGTATAGTTACATACGGTGACATTGACGCCGCGGGAAGAAAGCCGAACACGAC[G/ A]GAGGAGGCTGAGGGAGGAGCAGTTGGAGTGGTAGAAAGTGAGGAGGTCTGCGAGGTTG GGAATTTTCAGCGTACTTGTAAATAAGGAGCAGCCGTACATTTCTTCGGACATAGTAGGGT TTAGGGTTTATGCTCGCCGGTGCTGTCACTGCGGCTGCAGGGTTGCGCTATTCAACAGCG T
>Gm20_32736480	GTGACATTGACGCCGCGGGAAGAAAGCCGAACACGACCGAGGAGGCTGAGGGAGGAGCA GTTGGAGTGGTAGAAAAGTGAGGAGGTCGTCGAGGTTGGGAATTTTCAGCGTACTTGTAAATA AGGAGCACGCCGTACATTTCTTCGGACATAGTAGGGTTTAGGGTTTATGCTCGCCGGTG[C/ T]TGTCACTGCGGCTGCAGGGTTGCGCTATTCAACAGCGTAGGGTGGGGCTGGGCTTATGT CCGGTTGCCCATATCTTTTTGTTTTCTTTTTCTTTTTCTTTTTGCTTATGTCGGATAAATAA AAATTATTTTATTAGTATAAAAAATATAATAATTTTATCAAATCATTAAAGTTTGTGG
>Gm20_32736502	AAAGCCGAACACGACCGAGGAGGCTGAGGGAGGAGCAGTTGGAGTGGTAGAAAAGTGAGG AGGTGCTCGAGGTTGGGAATTTTCAGCGTACTTGTAAATAAGGAGCAGCCGTACATTTCTT CGGACATAGTAGGGTTTAGGGTTTATGCTCGCCGGTGCTGTCACTGCGGCTGCAGGGTT[A /G]CGCTATTCAACAGCGTAGGGTGGGGCTGGGCTTATGTCCGGTTGCCCATATCTTTTTG TTTTCTTTTTCTTTTTCTTTTTGCTTATGTCGGATAAATAAAAAATTATTTTATTAGTATAAAAA ATATAATAATTTTATCAAATCATTAAAGTTTGTGGGATTATTTGAATAATGCTCAA
>Gm20_32766243	GCGGCGGAGAGCCTCCACGAGGCGGCGGAGTAGATCCGCTGCTCCTCTTCGGATCTCC ACGGGGAGGCGGCGAGCCGAATTTTATCGGCGGCGCGTGGTCTCCGATTTTCTGCGCT CTTGTGATTGGATTCTCGCAACGTGTGCGGAATTACTAGAATCTACGCTCGGGTCGAAAG[G /A]CGCCATAGATGATTGCAATTGAATGATTTGTTCCGGTGAGATGAAGAGAGAAGAAATTGA ATGAACAATGAAGAACGAAAGAAGAAGTTGATGGTGTGTTGAAGCCAAGTGAGGGGAAGAA AATAGAAGAAAAGAAAGAGTTTTAAAGGAGTCAAAGTCACGGGGAACACGCGTCTCTTA
>Gm20_32766318	CGAATTTTGATCGGCGGCGGCGTGGTCTCCGATTTTCTGCGCTTCTTGTGATTGGATTCT CGCAACGTGTGCGAATTACTAGAATCTACGCTCGGGTCGAAAGACGCCATAGATGATTGCA ATTGAATGATTTGTTCCGGTGAGATGAAGAGAGAAGAAATTGAATGAACAATGAAGAA[G/A]G AAAGAAGAAGTTGATGGTGTGTTGAAGCCAAGTGAGGGGAAGAAAATAGAAGAAAAGAAAG AGTTTTAAAGGAGTCAAAGTCACGGGGAACACGCGTCTTTAGTTACTCACATTTATTTT GCGCCTACTGTTTTCTTCCGCTCGTTTTTCTGTTTTTTTATCTATTAATAATAATA
>Gm20_36069702	CTCAACAACCTCAGCACAAATCCTCATTCCCAACATTTGTTGCTCCAAGTTGTTATTTTCTG AGATGGTGGAAGGTTTGAATGCATCTTAGGAGAAAGGGTAGCAAGGAGACCATGGACAA CAAAGAGGATGGTGTCTTAAGCTCAGGGGATGTAAATTCGGAAAGCTGAGCTACCT[A/T] CTCCATAGCCACACCCAACTAAATTACTAATATAAATTGAAAATAAACTGAAAATATAGTTAA ATGAAATAGGCATGCACAAGTTAATAATGTCGCAATACAATTACGAAGGAATACAAGAGAA CATGCTTCTTTAAATGTTTGAATGAATACTTGGATTATCAAATCCTCAAACCTC
>Gm20_39942636	TTATTACTTTTTCTTAAGTTCTATAATTCAGTATTTACTTATTAATCATCCGTGCTCTGACCTT ATAGTTTAAAGGAACCTTTTTGCCTTATTTTTACATTATATAGGTAGCAAAAAAATGATT GGATAAAGAAAGAAGCAAGTGGAAGTTTAAATTTATTGGATAAATATCTCT[C/A]AGGTTTA GAGAATTGAAAAGATTTTCATTGATAATTGAGCAACACAATGAGTGCTCCACCTCATCATGA TGATGACGACGACAACGAAGAAGTTTTCTTGATGAGTCTGACATCATCCATGAAGTTGCC ATGGATAATGAAGATCTTCCTGACGCTGATGATGATTCTGAACCTCCTCGG
>Gm20_41042731	CATTTGTTAGTGATGCTGAGCAGCTGTGCTTGGTGTGTCACATTAGCTTGTTCAGGAAAT TTTGACAGTGGTATTTCAATTTTTATCATGCAAATCTTTATAGGACACCATGTTCAATGTCCT GGAATTTGGTCTTCCATGATTTGGCTTGGCATATTTTCACTTCAGAAATCAATA[G/T]ATTCC CTCAATACTGTATCTAGTTGAAGATAATTAGTAAGTCTGAGTTATACATCTCAAACACACTTG TTACTAAAGGTACCGCTTATTGTGATGGTGTACAAGTGTTAATGAGATTTCAATTTGATATGTT AACTTCGCAACCGTACCGCTTATATTCTAGGAGTTATTGTATTCTGTGCA
>Gm20_42627040	TGACCCCTTATAGCTTATTATCTGTTATTTCTCAGAACTGCGTTTGAGAATCCATTACCAGTC CTATCAAGTATAATCAACTCATTTCTCATTTGCAAATTGGTAGTATTTTAAACCATGCTATATA ACCTAAGAGTCAAGGACATTGACGCCAGAAATTTGGGGAAAGAAAAAACACAA[A/C]AAAAA AGAAGCAAACTTAAAAAGAAGGCATACTTCCTCCTCGTGAAGGGATGTTACTGTTTGGTC



	TCCGCTCTTCAATGTATACGTGTCTTCTGCTACCTCAACAGATCCTGCCTGCAAAGATACA AAACAAACATGAGCCAACAGATTCTGAAATCAGAAAAAAGAGACTTGTAAG
>Gm20_ 4262704 3	CCCCTTATAGCTTATTATCTGTTATTTCTCAGAACTGCGTTTGAGAATCCATTACCAGTCCTA TCAAGTATAATCAACTCATTCTCATTTGCAAATTGGTAGTATTTTAACCATGCTATATAACC TAAGAGTCAAGGACATTGACGCCAGAAATTTTGGGGAAGAAAAAACACAATAA[T/C]AAAGA AGCAAAACTTAAAAAGAAGGCATACTTCCTCCTCGTGAAGGGATGTTACTGTTTGGTCTCC GCTCTTCAATGTATACGTGTCTTCTGCTACCTCAACAGATCCTGCCTGCAAAGATACAAAA CAAACATGAGCCAACAGATTCTGAAATCAGAAAAAAGAGACTTGTAAGGTG
>Gm20_ 4267063 4	GGGTTTCTGCGTCCAAGAGAATCAAGGCTCTTTGACTCTGGGTTGGCATAGCCGCCCCCT AATTGCAGCTTCGGCTTGGCAAGCCGCGCCTCTGGGAGACGACGAAACGCTGCGTTTTGA GCATTGAGGAAAAAGAACAGAACAGACAGATCACTTGTTTTGTTTATTCTAGTAACTA[G/T] TAGCAGTAGTAGTACACGGGAAGAGTGGTGAATAATAAATGTGGTCATTTTTGCGTAAAA TTCTTCTGTTGTTGAATTCTCTATCACACGTGACTTTGGTTGGGGTTTTCTCTTTTGTCAA GAACCAACCAAGTGCTTTTCTTTTTTAGTCAATTGAGTCATACCATTGCCT
>Gm20_ 4269046 1	TTCCACACGCGAGGAAAGGTTACATAACAAGAACGTATGCTAGACAGTAAACTCTTTATTA TACTGATCAATCCCAAGGTTTTGTTTTCGAACATTACACCACTAAGATGTCTCAGAGAAACG ACACATAACGTAACATAACATAACACTATATAGTACTACATAATCCAATTTCAA[C/T]CCATC TTGGCATCACTAGTAGCATGCATTTTAGCACATTAATCAAATAAGCTCCCATCAAAGCATCT TCTAACAGCTTCTTTAGAGAGGAAACCTTCCTCATCCCTAGCCAGAATGTACAGAACCCCC CATTCAAGTTTAGCTGCAAGCCTGCAAATTAGTAAAACACACATTCCATATA
>Gm20_ 4269054 6	TTTTCGAACATTACACCACTAAGATGTCTCAGAGAAACGACACATAACGTAACATAACATAC ACACTATATAGTACTACATAATCCAATTTCAACCCATCTTGGCATCACTAGTAGCATGCATTT TAGCACATTAATCAAATAAGCTCCCATCAAAGCATCTTCTAACAGCTTCTTTAG[C/A]GAGGA AACCTTCTCATCCCTAGCCAGAATGTACAGAACCCCCATTCAAGTTTAGCTGCAAGCCT GCAAATTAGTAAACACACATTCATATATGTAACATTTTCATTGGATCTATACAATGAAAGTA ATTTGAGATGTGAGAAAATTCGCCAGCCAAATATGTCAAACGCATTTCCGT
>Gm20_ 4446496 5	GAATTAAGAAAAATAATTCCTAAAAGACTGTTAGTTCCAGGCTCAAATTCATATCAGACTA AAGGATACCTATATTTCAAGCTTGTGCTGGCTAAGAATGTAGTTTTCTTATTTTCTTAAGTT GAGAGTATTATGCCATGGACTTCACCACCAAGAAGCAAACTAAAAATTAGCAT[G/A]CCTA TCTTTATAAGCCGAAGCCAAATTAGCATGGGCTTCAGCCATGGTGGGTCTGACAGCAATTG CCCGTATGTAATCCTGGATAGCATCACTAACTCTACCAATCTCCTTATACGTATTGCCTCTA TTAACTAGCCCATCAGCTGCCAAGGGATCAATGCGGAGGACCTCATTGTAGCA
>Gm20_ 4465701 9	AATTCCTCGAACAGACCAGGACTCCGCATGGAATCATCTAAAACGTGTGTGACAGTTACCAG CCAAAAGTCTAGCTTTCTTGTGGTCACTTTCCAGCTTCCCAACATTTTCTGCAAGGCAACA AGTGCAGCGGCACAAAGGATGCCAATCTGTCTCATTCCACCTCCTAAGGTTTTCCG[A/G]AG TCGTCTAGCCTAAAAAATAATAAACTAGACTTCAACATCAGTTGCTCAGGAAGTAACATA CCGTCTATTAATTTCCCTGTAAAATTATTGAACCTACCTTGGCAATAAAATTTTGGAAACCAAC AATAACAGATCCAACCTGGAGCACCTATACCTTTAGATAGGCAAACCTAATCAT
>Gm20_ 4510435 5	AACAAGGTGCTACCTGCTTTGTGTAAGATCCAGCAAGAAAGAAATTCCTCACTGGCGTCTTT TGATCAGGTCTATATGGATCTTTACCAGGTCCCTCACGGTACAAAGATTGACCAATTTTAAC AACAGAAGACCAAGTGAATTCCTTCCAAACCTTGAGATGATGGGAACAGCGCTAAAACC[T/C]AAT AAAATTTTGAATCAATGTAATGACAAGAGAAGTGAATATCTACCATTGATTAACAAAAACCC AAATAGACTACAAAATGGAACCTGTGATTTGATTTGTGATGGAAATTATACAAAATGCTATG CTGATGATATCAGAACAAATCCAGTCAGCCATCAAATTAGCAGCATGAACAAA
>Gm20_ 4575181 7	TTTGTGAGTGTTACTGTTGAATCGATTGGATTGGCATTAGGTTGATCTGAAATACGCAATTG ACATGGCGAGCGCGATGGTAGAGGACGCGAAGTTCGAAGATGACCAACTAGCCAATATGA CCACCGACGACATCGTCAGAGCTTCACGTCTTCTCGACAACGAGATTTCGCATCCTCA[C/T]G GTATACCTTTTTCTTCTGTTTCTTTTTCCCTGCTTCTGTAACCTAACCCCTATCGATTA CACTATCTTTTTCTTCTTCCGTTTTCTTAGCCATTAGGGTTTTCTGATTTAATTGAATTGTC CTTCGTGACTGAAGGAAGAGTTGCAGAGGACCAATTTGGAATTGGAATCGTA
>Gm20_ 4611685 4	ATTTATAAAACATTGACCATATAAGTCAGGCAAATATGCACCTAAGAGTTTCATATCCTGGA GGAGTCATGGCAGGAAAATGCCACATGCAAGTTGCCAATTGCAAATCACAGGTAATATACT CACCCAAACATCTCCAGGAGAAGGGAAACCTCTTCTCATTAGGAGCTTGAATAAT[T/A]TG AGGAATAGAAAAGAGTTAACCTGCGGTTAACCTCCATTATCAGCAAAAACAAACAAATTGA CTCATCCAGAATGATACTCTTTTACCAATATAATGATAATAGTAGAACATAAAATTGCATACA CCAAAGAAAACCAAGGAACAAGAATGTATGTTCACTAGATCAGTATAGATT

**Supplemental File S1.** This file contains small scripts required to run SNPscript.R I have found many of these functions are also useful for other analyses.

```
#This script was written in the R programming language to determine intervals of
significant SNP density based on a bootstrap method.
#Required input for the script is a list of SNPs with chromosomal positions.
Output of the script is similar to Figure 2 and Figure 3.
#Created on 04/06/10 by Andrew Severin andrewseverin@gmail.com
#Iowa State University
#####
#This function will give the index number in a matrix given the rowname.
IndexFromGeneCall<-function(inputmatrix,rowname){
  match(rowname,rownames(inputmatrix),nomatch=0)
}
#####
#plotting function
#this function will generate intervals of significant clustering of SNPs on soybean
chromosomes scaled to the longest chromosome.
ChromosomePlot<-
function(chromosomelength,intervalstart,intervalend,intervalheight,maxNumGene
sInCluster,maxchromosomelength,binScales,currentBinScale,SNPcoords){
  #The axis labels require resizing depending on the width of the plot
  #the following is an estimate of the resizing required
  if (maxchromosomelength<750000){
    XcexVar<-1
  }
  if (maxchromosomelength>750000 & maxchromosomelength<12000000){
    XcexVar<-(-0.3125)*log(maxchromosomelength/1000000)+0.91
  }
  if (maxchromosomelength>12000000){
    XcexVar<-0.1
  }
  #I make use of the rect function that has input as (xleft, ybottom, xright, ytop)
  #keep in mind everything is scaled to Gm18, the largest chromosome
  xleft=(1/maxchromosomelength)*intervalstart
  ybottom=0
  width=(intervalend-intervalstart)/maxchromosomelength
  averagepos<-(intervalend+intervalstart)/(2*maxchromosomelength)
  average<-(intervalend+intervalstart)/(2)
  xright=xleft+width
  ytop=intervalheight/maxNumGenesInCluster
  rect(0,0,chromosomelength/maxchromosomelength,-0.04) #this draws the
chromosome on the bottom
```

```

rect(SNPcoords/maxchromosomelength,0,SNPcoords/maxchromosomelength ,-
0.04,lwd=.1) #this draws the location of each SNP
rect(xleft,ybottom,xright,ytop,border="black",col=rainbow(length(binscales))[whic
h(binscales==currentBinScale)]) #Significant Intervals of SNPs
text(chromosomelength/(2*maxchromosomelength),-
.02,labels=paste("chromosome",i),cex=.5) #chromosome name
text(SNPcoords/maxchromosomelength-2000/maxchromosomelength,-
.02,labels=rownames(SNPcoords),cex=XcexVar,srt=90) #location of each
Interval
axis(1,tick=T,at=intervalstart/maxchromosomelength,labels=intervalstart,cex.axis
=XcexVar,las = 2,lwd=.5) #axis 1
axis(2,tick=T,at=seq(0,1,1/maxNumGenesInCluster),labels=0:maxNumGenesInC
luster,cex.axis=.8) #axis 2
ablineMulti<-function(i){abline(i,0,col="darkgrey",lwd=.1)} #Creates a grid at 2
SNP intervals
sapply(seq(0,1,2/maxNumGenesInCluster),ablineMulti) #sapply to create the grid
}
#####
#this function is not used in the SNPscript but is a handy little function.
#Used in a similar script for clustering genes.
identifyGenesOnChromosomeForSoybean<-function(GeneList){
#this loop identifies all gene model names (glymas) on a specified chromosome
if (i<10){
glymas<-GeneList[grexp(paste("0",i,"g",sep=""),GeneList)]
}else{
glymas<-GeneList[grexp(paste(i,"g",sep=""),GeneList)]
}
return(glymas)
}
identifySNPSONChromosomeForSoybean<-
function(snpList,chromosomeNumber){
#this loop identifies all SNPs on a specified chromosome
if (chromosomeNumber<10){
snps<-
snpList[grexp(paste("Gm0",chromosomeNumber,sep=""),rownames(snpList)),]
}else{
snps<-
snpList[grexp(paste("Gm",chromosomeNumber,sep=""),rownames(snpList)),]
}
return(snps) #this will return the snp matrix that corresponds only to the
chromosome of interest
}
#####
#this section will take the same number of genes and simulate how the genes will

```

```

fall into the bins based on the 1000(or numofsims) random collections of genes
simulateData<-
function(numofsims,AllGeneCalls,geneCoordinates,SNPS,chromosomeNumber){
#matrix for storing simulations
genesAll<-identifyGenesOnChromosomeForSoybean(AllGeneCalls)
genesSample<-sample(genesAll,dim(SNPS)[1]*numofsims,replace=T)
AllindexCoords<-
function(i){IndexFromGeneCall(geneCoordinates,genesSample[i])}
sampleIndex<-sapply(1:length(genesSample),AllindexCoords)
genesSample<-matrix(genesSample,ncol=dim(SNPS)[1])
sampleIndex<-matrix(sampleIndex,ncol=dim(SNPS)[1])
return(list(genesSample=genesSample,sampleIndex=sampleIndex))
}
#####
#Boostrap function for clustering on a chromosome
#generation of the bin sizes across the chromosome
clusterByBoostrap<-
function(chromosomelength,binsize,geneIndexValues,SNPCoordinates,AllGeneC
alls,numofsims,bootData){
print(SNPCoordinates)
numBins<-floor(chromosomelength/binsize)
#this section will calcluate how many of of the SNPs we are interested in fall into
each bin
breaks1<-seq(0,chromosomelength,binsize)
chromBinsFind<-findInterval(SNPCoordinates,breaks1, rightmost.closed=T)
chromBins<-hist(chromBinsFind,breaks=seq(0,length(breaks1),1),plot=F)$counts
print(chromBins)
sampleIndex<-bootData$sampleIndex #actual data
chromBinsSample<-matrix(0,numofsims,length(breaks1)) #simulated data
for (j in 1:numofsims){
#generation of the bin sizes across the chromosome for the simulated data
breaks1<-seq(0,chromosomelength,binsize)
chromBinsSam<-findInterval(geneCoordinatesAve[sampleIndex[j,]],breaks1,
rightmost.closed=T)
chromBinsSample[j,]<-
hist(chromBinsSam,breaks=seq(0,length(breaks1),1),plot=F)$counts
}
#Average and standard deviation of the simulated data
chrombinsAve<-colMeans(chromBinsSample)
chrombinsSD<-sd(chromBinsSample)
#this section is the determination of the bins that are significant
over3stdev<-which((chromBins-(chrombinsAve+3*chrombinsSD))>0)
over3stdevBy<-(chromBins-(chrombinsAve+3*chrombinsSD))[over3stdev]
over3stdevZscore<-round(((chromBins-

```

```

(chrombinsAve))[over3stdev])/chrombinsSD[over3stdev],2)
#this if statement is required in case no intervals are found to be significant
if (length(over3stdev)==0){
print("No significant intervals found")
return(0)
}else{
significantIntervals<-
matrix(c(breaks1[over3stdev],breaks1[over3stdev]+binsize),length(over3stdev),2)
}
#append number of genes in the bin and the zscore to significantIntervals
significantIntervals<-
matrix(cbind(significantIntervals,chromBins[over3stdev],over3stdevZscore),ncol=
4)
significantIntervals<-
matrix(significantIntervals[sort(significantIntervals[,1],index.return=T)$ix,],ncol=4)
significantIntervals<-
matrix(significantIntervals[which(significantIntervals[,3]>1),],ncol=4)
print(significantIntervals)
if(dim(significantIntervals)[1]==0){
return(0)}else{
return(significantIntervals)
}
}
}

```

#This script is used to cluster SNPs onto Soybean chromosomes. Requires  
SNPsource.R and .RDataSNP

#This script was written in the R programming language to determine intervals of  
significant SNP density based on a bootstrap method.

#Required input for the script is a list of SNPs with chromosomal positions.

Output of the script is similar to Figure 2 and Figure 3.

#Created on 04/06/10 by Andrew Severin andrewseverin@gmail.com

#Iowa State University

#required libraries

library(gplots)

source('SNPsource.R')

load(".RDataSNP")

#starting parameters

dir<-"./"

numofsims<-3

SNPsofInterest<-read.table('./exampleSNPsFile.txt') #list with SNPs of interest

numberOfChromosome<-20 #this variable will allow you to loop through the first  
X chromosomes (see for loop below)

#this section is optional if you would like to have multiple bin sizes uncomment

#StartingBinsize<-6000000 #important the the vector in this forloop results in

```

binsizes that include the binsizes before it
#binscales<-c(1,2,6,12,60,120) #for binsize 6M 3M 1M 500K 100K 50k
#For multiple bin sizes comment out this block of code
StartingBinsize<-500000 #Here I chose just one binsize
binscales<-c(1)
#variables calculated from the input parameters
geneCoordinatesAve<-
matrix(round((geneCoordinates[,3]+geneCoordinates[,4])/2),ncol=1)
rownames(geneCoordinatesAve)<-rownames(geneCoordinates)
chromosomelengthAll<-chrom[,4]
maxchromosomelength<-max(chrom[,4])
significantIntervalsOrig<-0
#this for loop will cycle through each chromosomes.
for (i in 1:numberOfChromosome){
#for (i in numberOfChromosome:numberOfChromosome){ #This line can be
uncommented if you want to run it on a specific chromosome
dir.create(paste("./",i,sep="")) #create directory to export outfiles
chromosomelength<-chromosomelengthAll[i]
maxNumGenesInCluster<-0 #initiate a variable that will be needed later for
plotting
SNPs<-identifySNPSONChromosomeForSoybean(SNPsofInterest,i) #this
function will identify the SNPs on each chromosome as it goes through the loop
if (dim(SNPs)[1]<3){ #No need to look at chromosomes that do not have at least
3 SNPs
print(SNPs)
next()
}
bootData<-simulateData(numofsims,AllGeneCalls,geneCoordinates,SNPs,i)
#generate the simulated data (See SNPsource for code)
#binSize (for loop) will cycle through the binsizes determined above
for (b in binscales){
binsize<-StartingBinsize/b
print(binsize)
appendtofilename<-paste("_",binsize/binscales,sep="") #this variable is used for
the outputfiles to distinguish between bins
SNPcoords<-SNPs[,1] #for retrieval of the coordinates of the SNPs of interest
#function to do bootstrap method
significantIntervals<-
clusterByBootstrap(chromosomelength,binsize,geneCoordinatesAve,SNPcoords,
AllGeneCalls,numofsims,bootData)
print(significantIntervals)
if (b==binscales[1]){
#open a pdf file
pdf(file=paste(i,"/chrom",i,"ALL",".pdf",sep=""),paper="special",height=7,width=10

```

```

0)
plot(0:1, 0:1, type="n", axes=FALSE, ann=FALSE)
}
#if there are no significant Intervals go to the next binsize in the loop
if(significantIntervals==0){
next
}
#This block estimates the required Y dimension for plotting and works for most
cases
if (maxNumGenesInCluster==0){
maxNumGenesInCluster<-max(significantIntervals[,3])+1
}
#required input variables for the plotting function. ChromosomePlot can be found
in SNPsource.
intervalstart<-significantIntervals[,1]
intervalend<-significantIntervals[,2]
intervalheight<-significantIntervals[,3]
currentBinScale<-b
ChromosomePlot(chromosomelength,intervalstart,intervalend,intervalheight,max
NumGenesInCluster,maxchromosomelength, binScales,
currentBinScale,SNPcoords)
if(b==binscales[length(binscales)]){
#now that the plotting is finished, close the pdf file
dev.off()
}
#write to file gene lists with intervals that are significant
colnames(significantIntervals)<-
c('intervalstart','intervalend','numberInInterval','ZscoreaboveBootstrap')
write.table(significantIntervals,file=paste(i,"/clusterTable",i,appendtofilename,".txt"
,sep=""),append=T,quote=F,col.names=T)
}
#this commands save the R sessions for each chromosome into each
chromosome folder respectively.
save.image(file = paste(i,"/.RData",i,sep=""))
}

```

```

#the two columns are identical required to read in as a table in the script
position position
Gm01_53617124 53617124 53617124
Gm02_5687687 5687687 5687687

```

Gm02\_42350182 42350182 42350182  
Gm03\_36460374 36460374 36460374  
Gm03\_36554101 36554101 36554101  
Gm03\_36559857 36559857 36559857  
Gm03\_36559926 36559926 36559926  
Gm03\_36560002 36560002 36560002  
Gm03\_36952394 36952394 36952394  
Gm03\_36959955 36959955 36959955  
Gm03\_36996777 36996777 36996777  
Gm03\_36997184 36997184 36997184  
Gm03\_36997185 36997185 36997185  
Gm03\_37024958 37024958 37024958  
Gm03\_37027198 37027198 37027198  
Gm03\_37144398 37144398 37144398  
Gm03\_37165409 37165409 37165409  
Gm03\_37827399 37827399 37827399  
Gm03\_37828684 37828684 37828684  
Gm03\_37828791 37828791 37828791  
Gm03\_37832228 37832228 37832228  
Gm03\_37863675 37863675 37863675  
Gm03\_38065066 38065066 38065066  
Gm03\_38083417 38083417 38083417  
Gm03\_38117453 38117453 38117453  
Gm03\_38117485 38117485 38117485  
Gm03\_38132996 38132996 38132996  
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